Clone ID		ExperimentID HumanRNAAccession HumanProtAcces LocusLink	HumanProtAcces	LocusLink	Gene Symbol	Gene Description
cr0471	4C	NM_144981	NP_659418	196294	FLJ25059	hypothetical protein FLJ25059
cr0682	4C	NM_001030	NP_001021	6232	RPS27	ribosomal protein S27 (metallopanstimulin 1)
fcr0018	4C	NM_016208	NP_898880	51160	VPS28	vacuolar protein sorting 28 (yeast)
fcr0056	4C	NM_007112	NP_009043	7059	THBS3	thrombospondin 3
fcr0061	4C	005119	NP_005110	9967	THRAP3	thyroid hormone receptor associated protein 3
fcr0139	4C	NM_032142				
fcr0224	4C	NM_003801	NP_003792	8733	GPAA1	GPAAIP anchor attachment protein 1 homolog (yeast)
fcr0253	4C	XM_098004				
fcr0272	4C	NM_006233	NP_006224	5438	POLR2I	polymerase (RNA) II (DNA directed) polypeptide I,
	4C	NM_001636	NP_001627	293	SLC25A6	solute carrier family 25 (mitochondrial carrier; a
fcr0593	4C	NM_014780	NP_055595	9820	CUL7	cullin 7
fcr0608	4C	NM_014052				
fcr0707	4C	NM_005770	NP_005761	10169	SERF2	small EDRK-rich factor 2
fcr0712	4C	NM_000559	NP_000550	3047	HBG1	hemoglobin, gamma A
	4C	NM_032378	NP_115754	1936	EEFID	eukaryotic translation elongation factor 1 delta (
fcr0776	4C	NM_032268	NP_115644	84937	ZNRF1	zinc and ring finger protein 1
fcr0779	4C	NM_002882	NP_002873	5902	RANBPI	RAN binding protein 1
fcr0824	4C	NM_000182	NP_000173	3030	HADHA	hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Co
	4C	NM_030926	NP_112188	81618	ITM2C	integral membrane protein 2C
	4C	NM_021962	NP_068781	29	ABR	active BCR-related gene
	4C	NM_001416	NP_001407	1973	EIF4A1	eukaryotic translation initiation factor 4A, isofo
fcr0955	4C	NM_006886	NP_008817	514	ATP5E	ATP synthase, H+ transporting, mitochondrial F1 co
				7140	TNNT3	troponin T3, skeletal, fast
107990	40	NM_005833	NP 005824	10244	RAB9P40	Rah9 effector n40

fcr0997	4C	NM 005620	NP 005611	(808)	1100013	0100
fcr0999	4C	977700 MIN	NID 000770	0202	STOOATT	\$100 calcium binding protein A11 (calgizzarin)
fcr1068	40	1700 MIN	NP_002/69	5660	PSAP	prosaposin (variant Gaucher disease and variant me
fcr1173	46	NIM 004242	NP_006256	5885	RAD21	RAD21 homolog (S. pombe)
6/11/2	+6	NM_004343	NP_004334	811	CALR	calreticulin
fcr1305	4C	NM_014892				
fcr1337	4C	NM_004145	NP_004136	4650	МҮО9В	myosin IXB
fcr1463	4C	NM_001918	NP_001909	1629	DBT	dihydrolipoamide branched chain transactions (E2)
fcr1557	4C	NM_022356	NP_071751	64175	LEPRE1	leucine proline-enriched proteoglycan (learness)
fcr1633	4C	NM_014047	NP_054766	28974	HSPC023	HSPC023 protein
fcr1756	4C	NM_005327	NP 005318	3033	ОЗНОРН	
fcr1833	4C	NM_005343				A dehydrogenase, short ch
fcr1844	4C	NM_003492	NP 003483	8269	CXorf12	chromosom V
fcr1855	4C	NM_001866	NP_001857	1349	COX7B	Cytochrome Covidees on house Milli
fcr1879	4C	NM_003428	NP_003419	7637	ZNF84	zinc finger protein 84 (HPF2)
	4C	NM_033251	NP_150254	6137	RPL13	ribosomal protein L13
	4C	NM_017459	NP_059453	4237	MFAP2	microfibrillar-associated protein 2
	40	NM_014720	NP_055535	9748	SLK	SNF1 sucrose nonfermenting like kinase (veast)
	4C	NM_005252	NP_005243	2353	FOS	v-fos FBJ murine osteosarcoma viral oncogene homol
	4C	NM_014300	NP_055115	23478; 157708	SPC18	signal peptidase complex (18kD)
	4C	NM_002074	NP_002065	2782	GNB1	guanine nucleotide binding protein (G protein), he
	4C	NM_005898	NP_976240	4076	MHSI	membrane component, chromosome surface marker
	4C	NM_000972	NP_000963	6130; 147346	RPL7A	ribosomal protein L7a
ICr216/ 4	4C	NM_032860	NP_116249	84946	C6orf93	chromosome 6 open reading frame 93
fcr2196 4	4C	NM_014863				
fcr2276 4	4C	NM_007262	NP_009193	11315	PARK7	Parkinson disease (autosomal recessive, early onse
NY1-4308219v1	/1					4

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ICr2299	46	NM_000154	NP 000145	2584	GALK	09 9Ctokingse 1
fcr2573	4C	NM_000942	NP_000933	5479	PPIB	DeptidyInroly isomerase R (avolontilla D)
fcr2607	4C	NM_006758	NP_006749	7307	U2AF1	U2(RNU2) small nuclear RNA auxiliary factor 1
fcr2798	4C	NM_032830	NP_116219	84916	CIRHIA	cirrhosis, autosomal recessive IA (cirhin)
fcr2859	4C	NM_004223	NP_937826	9246	UBE2L6	
fcr2940	4C	NM_182470	NP_872271	5315; 400388	PKM2	pyruvate kinase, muscle
fcr2952	4C	NM_058246	NP_490647	10049; 387820	DNAJB6	DnaJ (Hsp40) homolog, subfamily B, member 6
tcr2958	4C	NM_019609	NP_062555	56265	CPXM	carboxypeptidase X (M14 family)
icr3053	40	NM_080425	NP_536351	2778	GNAS	GNAS complex locus
fcr3101	4C	NM_012073	NP_036205	22948	CCT5	chaperonin containing TCP1, subunit 5 (epsilon)
Icr3121	4C	NM_015934	NP_057018	51602	NOP5/NOP58	nucleolar protein NOP5/NOP58
fcr3155	4C	NM_003281	NP_003272	7135	IINNT	troponin I, skeletal, slow
Icr3181	4C	NM_005318	NP_005309	3005	HIF0	H1 histone family, member 0
ICF3323	40	NM_199330	NP_955364	9455	HOMER2	homer homolog 2 (Drosophila)
fcr3559	4C	NM_002615	NP_002606	5176	SERPINFI	serine (or cysteine) proteinase inhibitor, clade F
	4C	NM_005003	NP_004994	4706	NDUFABI	NADH dehydrogenase (ubiquinone) 1, alpha/beta subc
	4C	NM_002165	NP_851998	3397	IDI	inhibitor of DNA binding 1, dominant negative heli
	÷	NM_005507	NP_005498	1072	CFL1	cofilin 1 (non-muscle)
	4C	NM_005964	NP_005955	4628	МҮН10	myosin, heavy polypeptide 10, non-muscle
	4C	NM_016081	NP_057165	23022	KIAA0992	palladin
	4C	NM_006185	NP_006176	4926	NUMAI	nuclear mitotic apparatus protein
	4C	NM_006371	NP_006362	10491	CRTAP	cartilage associated protein
tcr4129	4C	NM_000466	NP_000457	5189	PEX1	peroxisome biogenesis factor
	4C	NM_004548	NP_004539	4716	NDUFB10	NADH dehydrogenase (ubiquinone) I beta subcomplex
ICF42/2	40	NM_032592	NP 115981	84680	PHACS	1-aminocyclopropane-1-carboxylate synthese

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52 NP_004003 1351 52 NP_758436 23509 76 NP_99827 6774 24 NP_998827 6774 19 NP_976316 23600 19 NP_006810 10963 10 NP_000261 4860 NP_0002607 5187 1 NP_002607 5187 1 NP_003365 7416 N NP_003365 7416 N NP_066949 4357 N NP_071767 64223 O NP_07962 468 A NP_087962 468 A NP_08938 11052 C NP_089980 54732 H NP_059980 54732 H	ferritin, light polypeptide		2312	NF_00013/			
52 NP_004003 1351 COX8A 52 NP_758436 23509 POFUT1 76 NP_998827 6774 STAT3 24 NP_976316 23600 AMACR 28 NP_919313 152006 RNF38 19 NP_006810 10963 STIP1 70 NP_002607 5187 PER1 6 NP_002607 5187 PER1 33 NP_808853 83658 DNCL2A 4 NP_003365 7416 VDAC1 1 NP_006808 10961 C12orf8 6 NP_006849 4357 MPST 7 NP_006825 10245 TIMM17B Inchestor 6 NP_008938 11052 CPSF6 C NP_008958 11073 TOPBP1 Inchestor NP_059980 \$4732 HSGP3402 C	gpzotz protein	11001 6010	2612	ND 000137	NM 000146	4C	fcr5339 /
52 NP_758436 23509 POFUT1 76 NP_998827 6774 STAT3 24 NP_976316 23600 AMACR 28 NP_919313 152006 RNF38 19 NP_000261 4860 NP 6 NP_002607 5187 HF1 6 NP_002607 5187 PER1 3 NP_808853 83658 DNCL2A 0 NP_001601 53 ACP2 1 NP_003365 7416 VDAC1 1 NP_006808 10961 C12orf8 5 NP_006825 10245 TIMM17B In NP_008938 11052 CPSF6 C NP_008958 11073 TOPBP1 6	COOKING PROCESS	HSCD2SI 2C	54732	NP 059980	NM_017510	4C	fcr5316 /
S2 NP_004003 1351 COX8A 52 NP_758436 23509 POFUT1 76 NP_998827 6774 STAT3 24 NP_976316 23600 AMACR 28 NP_919313 152006 RNF38 19 NP_000810 10963 STIP1 6 NP_000177 3075 HF1 6 NP_002607 5187 PER1 3 NP_808853 83658 DNCL2A 4 NP_003365 7416 VDAC1 1 NP_006808 10961 C12ort8 5 NP_006849 4357 MPST 1 NP_005825 10245 TIMM17B 1 NP_659003 53827 FXYD5 1 NP_008938 11052 CPSF6 c	topoisomerase (DNA) II binding protein	TOPBPI	11073	NP_008958	NM_007027	46	
NP_004063 1351 COX8A 82 NP_758436 23509 POFUT1 76 NP_998827 6774 STAT3 24 NP_976316 23600 AMACR 8 NP_919313 152006 RNF38 9 NP_006810 10963 STIP1 0 NP_000261 4860 NP 6 NP_002607 5187 PER1 3 NP_808853 83658 DNCL2A 0 NP_001601 53 ACP2 4 NP_003365 7416 VDAC1 NP_0066949 4357 MPST NPST NP_0071767 64223 GBL OR NP_005825 10245 TIMM17B DR NP_659003 53827 FXYD5 F	cleavage and polyadenylation specific factor 6, 68	CPSF6	11052	NP_008938	NM_007007	6 6	
NP_004063 NP_758436 NP_758436 NP_998827 NP_998827 NP_976316 NP_976316 NP_919313 NP_006810 NP_000261 NP_000261 NP_000177 NP_002607 NP_002607 NP_001601 NP_00365 NP_000432 NP_006808 NP_006808 NP_006808 NP_0068049 NP_005825 NP_005825 NP_877962 NP_T58436 ROPEUT1 NP_NOT1767 NP_NOT1767	FXYD domain containing ion transport regulator 5	FXYD5	53827	NP_659003	NM_014164	\$ E	
NP_004003 NP_758436 NP_998827 NP_998827 NP_998827 NP_998827 NP_976316 23600 NP_976316 23600 NP_919313 152006 RNF38 NP_006810 10963 STIP1 NP_000261 A860 NP NP_002607 5187 PER1 NP_003365 NP_003365 NP_003365 NP_006808 NP_006808 NP_006808 NP_006808 NP_006808 NP_006809 NP_005825 NP_005825 NSTAT3 AMACR AMACR NP	activating transcription factor 4 (tax-responsive	ATF4	468	NP_877962	NM_0016/5		
NP_004003 NP_758436 NP_758436 NP_998827 NP_998827 NP_998827 NP_998827 NP_998827 NP_998827 NP_998827 NP_900810 NP_000810 NP_000261 NP_000261 NP_000261 NP_0002607 NP_002607 NP_001601 NP_003365 NP_003365 NP_000432 NP_006808 NP_0068949 NP_0068949 NP_0068949 NP_071767 NP_071767 NP_071767 NP_006808 NP_071767 NP_006808 NP_071767 NP_006808 NP_071767 NP_006808 NP_071767 NP_006808 NP_071767 NP_006808 NP_071767	translocase of inner mitochondrial membrane 17 hom	TIMM17B	10245	NP_005825	INIM 001/22	40	
NP_004003 NP_758436 NP_758436 NP_998827 NP_998827 NP_998827 NP_998827 NP_976316 NP_976316 NP_919313 NP_006810 NP_000261 NP_000261 NP_000261 NP_0002607 S187 PERI NP_003365 NP_003365 NP_000432 NP_006808 NP_006808 NP_0066949 NP_0066949 NPST NP	G protein beta subunit-like	GBL	64223	NP_071767	NIM 005024	40	
NP_003365 NP_006808 NP_006810 NP_0003365 NP_0006808 NP_0006810 NP_0006808 NP_0006808 NP_0006808 NP_0006808 NP_006808	mercaptopyruvate sulfurtransferase	MPST	4357	NP_066949	OZITZO MIN	40	
NP_004003 1351 COX8A NP_758436 23509 POFUT1 NP_998827 6774 STAT3 NP_976316 23600 AMACR NP_976313 152006 RNF38 NP_000261 4860 NP NP_000177 3075 HF1 NP_002607 5187 PER1 NP_808853 83658 DNCL2A NP_003365 7416 VDAC1 NP_000432 5172 SLC26A4		C12orf8	10961	NP_006808	NM_00417	40	fcr5075
NP_004063 1351 COX8A NP_758436 23509 POFUT1 NP_998827 6774 STAT3 NP_998827 6774 STAT3 NP_976316 23600 AMACR NP_919313 152006 RNF38 NP_000261 4860 NP NP_000177 3075 HF1 NP_808853 83658 DNCL2A NP_001601 53 ACP2 NP_003365 7416 VDAC1		SLC26A4	5172	NP_000432	NM_000441		
NP_002607 5187 PERI NP_004063 1351 COX8A 2 NP_758436 23509 POFUT1 5 NP_998827 6774 STAT3 4 NP_976316 23600 AMACR NP_006810 10963 STIP1 NP_000261 4860 NP NP_000267 5187 PERI NP_808853 83658 DNCL2A NP_001601 53 ACP2	voltage-dependent anion channel 1	VDAC1	7416	NP_003365	NM 00311		for4065
NP_002607 5187 PER I NP_004063 1351 COX8A 22 NP_758436 23509 POFUT1 NP_998827 6774 STAT3 NP_998827 6774 STAT3 NP_919313 152006 RNF38 NP_006810 10963 STIP1 NP_000261 4860 NP NP_000267 5187 PER I NP_808853 83658 DNCL2A	acid phosphatase 2, lysosomal	ACP2	53	NP_001601	NIM 001010	AC	fcr4954
NP_002607 5187 PERI	dynein, cytoplasmic, light polypeptide 2A	DNCL2A	83658	NP_808853	NIM 001710	40	fcr4927
NP_000061 1351 COX8A NP_758436 23509 POFUT1 NP_998827 6774 STAT3 NP_976316 23600 AMACR NP_919313 152006 RNF38 NP_000810 10963 STIP1 NP_000261 4860 NP NP_000177 3075 HF1	period homolog 1 (Drosophila)	PERI	5187	NP_002607	MINI 014152	40	fcr4885
NP_998827 6774 STAT3 NP_998827 6774 STAT3 NP_976316 23600 AMACR NP_919313 152006 RNF38 NP_006810 10963 STIP1 NP_000261 4860 NP	H factor I (complement)	nr i	0070	- 000 (01)	719C00 WN	4C	fcr4846
NP_998827 6774 STAT3 NP_976316 23600 AMACR NP_919313 152006 RNF38 NP_006810 10963 STIP1 NP_000261 4860 NB	nucleoside phosphorylase	HE N	3075	NP 000177	NM_000186	4C	fcr4832
NP_758436 23509 POFUT1 NP_998827 6774 STAT3 NP_976316 23600 AMACR NP_919313 152006 RNF38 NP_006810 10963 STIP1		ND	4860	NP 000261	NM_000270	4C	fcr4831
NP_758436 23509 POFUT1 NP_998827 6774 STAT3 NP_976316 23600 AMACR NP_919313 152006 RNF38	stress-induced-phosphoprotoin 1 (11, 70, 7)	STIPI	10963	NP_006810	NM_006819	4C	tcr4795
NP_758436 23509 POFUT1 NP_998827 6774 STAT3 NP_976316 23600 AMACR	ring finger protein 38	RNF38	152006	NP_919313	NM_194328	-	1014/02
NP_758436 23509 POFUT1 NP_998827 6774 STAT3	alpha-methylacyl-CoA racemase	AMACR	23600	NP_976316	NM_014324	3 7	fcr/1707
NP_758436 23509 POFUT1 protein O-fu	signal transducer and activator of transcription 3	STAT3	6774	NP_998827	NM_1392/6	5 8	for4763
COX8A cytochrome	protein O-fucosyltransferase 1	POFUT1	23509	NP_758436	NM_013332	40	fcr4699
100000	cytochrome c oxidase subunit 8A	COX8A	1351	NP_004065	NM_0040/4	40	fcr4634
	WD repeat and SOCS box-containing	WSB1	26118	NP_599027	NIM 004074	30	fcr4622
	zinc finger protein 151 (pHZ-67)	ZNF151	7709	NP_003434	NM_003443	40	fcr4477

25 4C NM_001861 NP_001852 25 4C NM_001402 NP_001393 70 4C NM_001402 NP_001393 29 4C NM_000527 NP_000518 89 4C NM_0001246 NP_073143 9 4C NM_0001946 NP_073143 10 4C NM_0017801 NP_000271 10 4C NM_0017801 NP_000271 10 4C NM_007108 NP_978820 11 4C NM_006196 NP_078820 12 4C NM_006415 NP_08896 13 4C NM_006415 NP_0058632 14 4C NM_006415 NP_005430 14 4C NM_004875 NP_005430 14 4C NM_004875 NP_005960 14 NM_005969 NP_005960 14 NM_006572 NP_006563	solute carrier family 9 (sodium/hydrogen exchanger	SLC9A3R2	9351	NP_004776	141A) CO / +OO _141A)		
3-4 NM_001861 NP_001852 1327 COX4II 25 4C NM_001402 NP_001393 1915; EEF1AI 70 4C NM_001402 NP_001393 1915; EEF1AI 39 4C NM_000527 NP_000518 3949 LDLR 39 4C NM_001946 NP_002119 3146 HMGBI 8 4C NM_001980 NP_0002119 3146 HMGBI 8 4C NM_001980 NP_000271 54918 CKLFSF6 3 4C NM_01780I NP_060271 54918 CKLFSF6 3 4C NM_0173476 NP_597733 171017 ZNF384 5 4C NM_004844 NP_078820 79594 FLJ12875 8 4C NM_006196 NP_08832 29855 UBNI 5 4C NM_006485 NP_060581 55150; FLJ10490 6 4C NM_005439 NP_005430 8079 MLF2 <	guanine nucleotide binding protein (G protein), al	GNA13	10672		2/C000 TAIN		
3-4 C NM_001861 NP_001852 1327 COX411 25 4C NM_001402 NP_001393 1915; EEF1AI 70 4C NM_001402 NP_001393 1915; EEF1AI 39 4C NM_000327 NP_000518 3949 LDLR 8 4C NM_001946 NP_002119 3146 HMGBI 8 4C NM_000998 NP_000219 3146 HMGBI 8 4C NM_0017801 NP_060271 54918 CKLFSF6 3 4C NM_017801 NP_060271 54918 CKLFSF6 3 4C NM_017801 NP_060271 54918 CKLFSF6 3 4C NM_017801 NP_060271 54918 CKLFSF6 3 4C NM_0173476 NP_597733 171017 ZNF384 5 4C NM_006196 NP_06187 5093 PCBPI 5 4C NM_005433 NP_055832 29855 <td< td=""><td>nucleosome assembly protein 1-like 4</td><td>NAP1L4</td><td>4676; 55434</td><td>NP_005960</td><td>NIM 008573</td><td></td><td></td></td<>	nucleosome assembly protein 1-like 4	NAP1L4	4676; 55434	NP_005960	NIM 008573		
94 4C NM_001861 NP_001852 1327 COX4II 25 4C NM_001402 NP_001393 1915; EEF1AI 70 4C NM_001402 NP_001393 1915; EEF1AI 8 4C NM_000527 NP_000518 3949 LDLR 9 4C NM_000998 NP_002119 3146 HMGBI 10 4C NM_000998 NP_000989 6168 RPL37A 0 4C NM_0017801 NP_060271 54918 CKLFSF6 3 4C NM_013476 NP_597733 171017 ZNF384 5 4C NM_016936 NP_996896 6923 TCEB2 9 4C NM_016936 NP_908832 29855 UBNI 4 4C NM_003164 NP_003155 6811 STX5A 5 4C NM_006415 NP_906381 55150; FLJ10490 6 4C NM_006415 NP_906349 8079 MLF		TEX261	113419	NP_003183	NIM 002020	10 6	
94 4C NM_001861 NP_001852 1327 COX4II 25 4C NM_001402 NP_01393 1915; EEF1AI 70 4C NM_001402 NP_01393 1915; EEF1AI 8 4C NM_000527 NP_000518 3949 LDLR 9 4C NM_000998 NP_002119 3146 HMGBI 8 4C NM_000998 NP_003143 1848 DUSP6 5 4C NM_000998 NP_060271 54918 CKLFSF6 5 4C NM_017801 NP_060271 54918 CKLFSF6 8 4C NM_017801 NP_07820 79594 FLJ12875 8 4C NM_024544 NP_078820 79594 FLJ12875 8 4C NM_006196 NP_06187 5093 PCBP1 5 4C NM_006196 NP_065832 29855 UBN1 5 4C NM_006196 NP_060581 55150; FLJ10	polymerase (RNA) I polypeptide C, 30kDa	POLRIC	9533	NP_976035	NM 144583	40 7	
4 4C NM_001861 NP_001852 1327 COX4II 5 4C NM_001402 NP_001393 1915; EEF1AI 5 4C NM_001402 NP_001393 1915; EEF1AI 389411 389411 389411 389411 389411 4C NM_000527 NP_000518 3949 LDLR 4C NM_001946 NP_073143 1848 DUSP6 4C NM_001946 NP_073143 1848 DUSP6 4C NM_017801 NP_060271 54918 CKLFSF6 4C NM_0133476 NP_597733 171017 ZNF384 4C NM_007108 NP_96896 6923 TCEB2 4C NM_006196 NP_08632 29855 UBN1 PCBP1 4C NM_003164 NP_08632 29855 UBN1 UBN1 4C NM_006415 NP_086381 55150; FLJ10490 PCBP1 4C NM_006415 NP_0863430 8079	polymerase (RNA) II (DNA directed) polypeptide D	POLR2D	5433	NP_004796	NM 004805		
4 4C NM_001861 NP_001852 1327 COX411 5 4C NM_001402 NP_001393 1915; EEF1A1 0 4C NM_001402 NP_001393 1915; EEF1A1 389411 389411 389411 389411 389411 4C NM_000527 NP_000518 3949 LDLR 4C NM_001946 NP_073143 1848 DUSP6 4C NM_001980 NP_000989 6168 RPL37A 4C NM_017801 NP_000271 54918 CKLFSF6 4C NM_017344 NP_078820 79594 FLJ12875 4C NM_006196 NP_006187 5093 PCBP1 F 4C NM_016936 NP_058632 29855 UBN1 UBN1 4C NM_006415 NP_060581 55150; FLJ10490 N 4C NM_006415 NP_005430 8079 MLF2 N	myosin, light polypeptide 6, alkali, smooth muscle	MYL6	4637	NP_524149	NM_021019	7 7	
4 4C NM_001861 NP_001852 1327 COX411 5 4C NM_001402 NP_001393 1915; EEF1A1 0 4C NM_138501 NP_001393 1915; EEF1A1 389411 389411 389411 GPSN2 391676 NP_002128 391676 DLR 4C NM_000527 NP_000518 3949 LDLR 4C NM_001946 NP_002119 3146 HMGB1 4C NM_017801 NP_000989 6168 RPL37A 4C NM_017801 NP_597733 171017 ZNF384 4C NM_024544 NP_96896 6923 TCEB2 4C NM_006196 NP_96896 6923 TCEB2 TCEB2 4C NM_003164 NP_058632 29855 UBN1 PCBP1 F 4C NM_006415 NP_060581 55150; FLJ10490 P 4C NM_006415 NP_847894 10558 SPTLC1 SPT	myeloid leukemia factor 2	MLF2	8079	NP_005430	NM_005439	16	C10233
4 4C NM_001861 NP_001852 1327 COX411 5 4C NM_001402 NP_001393 1915; EEF1A1 0 4C NM_138501 NP_012510 9524; GPSN2 0 4C NM_000527 NP_000518 3949 LDLR 0 4C NM_001946 NP_073143 1848 DUSP6 4C NM_001980 NP_060271 54918 CKLFSF6 4C NM_0133476 NP_597733 171017 ZNF384 4C NM_024544 NP_078820 79594 FLJ12875 4C NM_006196 NP_08863 6923 TCEB2 TCEB2 4C NM_016936 NP_058632 29855 UBN1 NP_08153 4C NM_003164 NP_060581 55150; FLJ10490 NP_010490	serine palmitoyltransferase, long chain base subun	SPTLC1	10558	NP_847894	NM_006415	40	001010
4 POINT NP_001852 1327 COX4II 5 4C NM_001402 NP_001393 1915; EEF1AI 5 4C NM_138501 NP_001393 1915; EEF1AI 6 4C NM_000527 NP_000518 394911 LDLR 7 4C NM_000527 NP_000518 3949 LDLR 8 4C NM_000998 NP_002119 3146 HMGBI 8 4C NM_000998 NP_000989 6168 RPL37A 9 4C NM_017801 NP_060271 54918 CKLFSF6 4C NM_0133476 NP_597733 171017 ZNF384 4C NM_007108 NP_90896 6923 TCEB2 4C NM_006196 NP_008187 5093 PCBP1 4C NM_003164 NP_003155 6811 STX5A s	hypothetical protein FLJ10490	FLJ10490	55150; 64130	NP_060581	NM_022165	40	E 150
4 4C NM_001861 NP_001852 1327 COX4II 5 4C NM_001402 NP_001393 1915; EEF1AI 0 4C NM_138501 NP_012510 9524; GPSN2 9 4C NM_000527 NP_000518 3949 LDLR 9 4C NM_001946 NP_073143 1848 DUSP6 5 4C NM_017801 NP_060271 54918 CKLFSF6 6 4C NM_0133476 NP_597733 171017 ZNF384 4C NM_024544 NP_078820 79594 FLJ12875 1 4C NM_006196 NP_006187 5093 PCBP1 PCBP1 4C NM_016936 NP_058632 29855 UBN1 P	syntaxin 5A	STX5A	6811	NP_003155	NM_003164	4	10010
4 C NM_001861 NP_001852 1327 COX4II 5 4C NM_001402 NP_001393 1915; EEF1AI 0 4C NM_138501 NP_012510 9524; GPSN2 9 4C NM_000527 NP_000518 3949 LDLR 9 4C NM_001946 NP_002119 3146 HMGBI 8 4C NM_001946 NP_073143 1848 DUSP6 9 4C NM_001980 NP_060271 54918 CKLFSF6 4 C NM_0133476 NP_597733 171017 ZNF384 4 C NM_024544 NP_078820 79594 FLJ12875 4 C NM_006196 NP_996896 6923 TCEB2 TCEB2	ubinuclein I	UBNI	29855	NP_058632	NM_016936	ō t	6
4 C NM_001861 NP_001852 1327 COX411 5 4C NM_001402 NP_001393 1915; EEF1A1 0 4C NM_138501 NP_012510 9524; GPSN2 9 4C NM_000527 NP_000518 3949 LDLR 9 4C NM_001946 NP_073143 1848 DUSP6 4C NM_001980 NP_000989 6168 RPL37A 4C NM_017801 NP_060271 54918 CKLFSF6 4C NM_024544 NP_597733 171017 ZNF384 4C NM_007108 NP_996896 6923 TCEB2	poly(rC) binding protein 1	PCBPI	5093	NP_006187	NM_01636	AC #	fcr5880
4 C NM_001861 NP_001852 1327 COX411 5 4C NM_001402 NP_001393 1915; EEF1A1 0 4C NM_138501 NP_612510 9524; GPSN2 9 4C NM_000527 NP_000518 3949 LDLR 9 4C NM_002128 NP_073143 1848 HMGB1 8 4C NM_001946 NP_073143 1848 DUSP6 4C NM_0017801 NP_060271 54918 CKLFSF6 4C NM_133476 NP_597733 171017 ZNF384 4C NM_024544 NP_078820 79594 FLJ12875 FLJ12875	transcription elongation factor B (SIII), polypept	TCEB2	6923	NP_996896	NM_00/10%	7) t	fcr5770
4 C NM_001861 NP_001852 1327 COX411 5 4C NM_001402 NP_001393 1915; EEF1A1 0 4C NM_138501 NP_612510 9524; GPSN2 0 4C NM_000527 NP_000518 3949 LDLR 0 4C NM_002128 NP_002119 3146 HMGB1 8 4C NM_001946 NP_073143 1848 DUSP6 6 4C NM_000998 NP_000989 6168 RPL37A 4C NM_133476 NP_597733 171017 ZNF384	hypothetical protein FLJ12875	FLJ12875	79594	NP_078820	NM_02100	AC 7	fcr5758
4C NM_001861 NP_001852 1327 COX4II 4C NM_001402 NP_001393 1915; BEF1AI EEF1AI 4C NM_138501 NP_612510 9524; GPSN2 4C NM_000527 NP_000518 3949 LDLR 4C NM_001946 NP_073143 1848 DUSP6 4C NM_000998 NP_000989 6168 RPL37A 4C NM_017801 NP_060271 54918 CKLFSF6	zinc finger protein 384	ZNF384	171017	NP_597733	NM_133476	4C	fcr5755
4C NM_001861 NP_001852 1327 COX4II 4C NM_001402 NP_001393 1915; BEF1AI 389411 4C NM_138501 NP_612510 9524; GPSN2 4C NM_000527 NP_000518 3949 LDLR 4C NM_002128 NP_002119 3146 HMGBI 4C NM_001946 NP_073143 1848 DUSP6 4C NM_000998 NP_000989 6168 RPL37A	chemokine-like factor super family 6	CKLFSF6	54918	NP_060271	NM_017801	† C	fer5/20
4C NM_001861 NP_001852 1327 COX411 4C NM_001402 NP_001393 1915; BEF1A1 EEF1A1 4C NM_138501 NP_612510 9524; GPSN2 4C NM_000527 NP_000518 3949 LDLR 4C NM_002128 NP_002119 3146 HMGB1 4C NM_001946 NP_073143 1848 DUSP6	ribosomal protein L37a	RPL37A	6168	NP_000989	NM_000998	5	C 2000
4C NM_001861 NP_001852 1327 COX411 4C NM_001402 NP_001393 1915; EEF1A1 389411 SEF1A1 4C NM_138501 NP_612510 9524; GPSN2 4C NM_000527 NP_000518 3949 LDLR 4C NM_000528 NP_002119 3146 HMGB1	dual specificity phosphatase 6	DUSP6	1848	NP_073143	NM_001946	6 6	810CIDI
4C NM_001861 NP_001852 1327 COX4II 4C NM_001402 NP_001393 1915; EEF1AI 4C NM_138501 NP_612510 9524; GPSN2 4C NM_000527 NP_000518 3949 LDLR	high-mobility group box 1	HMGB1	3146	NP_002119	NM_002128	40	6.55710
4C NM_001861 NP_001852 1327 COX411 4C NM_001402 NP_001393 1915; EEF1A1 4C NM_138501 NP_612510 9524; GPSN2 391676	low density lipoprotein receptor (familial hyperch	LDLR	3949	NP_000518	NM_000527	45	E 5550
4C NM_001861 NP_001852 1327 COX411 4C NM_001402 NP_001393 1915; EEF1A1		GPSN2	9524; 391676	NP_612510	NM_138501	40 4	fcr5500
4C NM_001861 NP_001852 1327 COX411	eukaryotic translation elongation factor alpha	EEFIAI	1915; 389411	NP_001393	NM_001402	7 7	for\$470
	cytochrome c oxidase subunit IV isoform 1	COX4II	1327	NP_001852	NM_001861	À t	fcr5/125

TIAL1 TIAI cytotoxic granule-associated RNA binding prot GHITM growth hormone inducible transmembrane protein KIAA2010 KIAA2010 WBSCR1 Williams-Beuren syndrome chromosome region 1 CSNK2B casein kinase 2, beta polypeptide	1460	NP_001311	1001320		
0			CONTRACTOR I	•	
	7458	NP_114381	NM_022170	40	
	55671	NP_115949	NM 032120		1
	27069	NP_055209	NM_032500		- 1
	7073	NP_071728	NM_022333	40	ferb1337
HNRPA0 heterogeneous nuclear ribonucleoprotein A0	10949	NP_006796	NM_006805	4C	
	51669	NP_057211	NM_016127	4C	
TPM1 tropomyosin l (alpha)	7168	NP_000357	NM_000366	4C	1
FLJ13352 hypothetical protein FLJ13352	79644	NP_078868	NM_024592	4C	1
PCBP1 poly(rC) binding protein 1	5093	NP_006187	NM_006196	4C	1
FSTL1 follistatin-like	11167	NP_009016	NM_007085	4C	ICT/58/
MTX1 metaxin I	4580	NP_942584	NM_002455	6	101/419
517; 341412 ATP5G2 ATP synthase, H+ transporting, mitochondrial F0 co	517; 341	NP_005167	NM_005176	+	627/10
IGF2 insulin-like growth factor 2 (somatomedin A)	3481	NP_000603	NM_000612	40	for 7403
GNB2 guanine nucleotide binding protein (G protein) he	2783	NP_005264	NM_005273	4C	fcr /042
COPB coatomer protein complex, subunit beta	1315	NP_057535	NM_016451	4C	fcr/026
MDK midkine (neurite growth-promoting factor 2)	4192	NP_002382	NM_002391	4C	fcr/004
P4HB procollagen-proline, 2-oxoglutarate 4-dioxyogenase	5034	NP_000909	816000 WN	4C	tcr6937
NAG	51594	NP_056993	NM_015909	4C	tcr6931
CCNB1 cyclin B1	891	NP_114172	NM_031966	4C	Icro881
KIAA0090	23065	NP_055862	NM_015047	4C	Icr6616
TIRAP toll-interleuk	114609	NP_683708	NM_052887	4C	fcr6592
RHOB ras homolog gene family, member R	388	NP_004031	NM_004040	4C	fcr6577
RBBP7 retinoblastoma binding protein 7	5931	NP_002884	NM_002893	4C	tcr6573
NUP54 nucleoporin 54kDa	53371	NP_059122	NM_017426	†c	icr6534

					1	1701 C801 LAIV
hypothetical protein FLJ12442	FLJ12442 h	64943	NP_075059	NM_022908	4C	fcrb2090 4
fibulin 1	FBLN1 fi	2192	NP_006478	NM_006486	4C	1
melanoma antigen, family D, 1	MAGEDI n	9500	NP_008917	NM_006986		
prostaglandin F2 receptor negative regulator	PTGFRN p	5738	NP_065173	NM_020440		
protein tyrosine phosphatase, non-receptor type 23	PTPN23 p	25930	NP_056281	NM_015466		
insulin-like growth factor 2 (somatomedin A)	IGF2 ii	3481	NP_000603	NM_000612		1
hemoglobin, alpha 2	HBA2 h	3040; 3039	NP_000508	NM_000558	40	
tubulin, gamma complex associated protein 2	TUBGCP2 t	10844	NP_006650	NM_006659	4C	1
myosin, light polypeptide 6, alkali, smooth muscle	MYL6 r	4637	NP_524149	NM_079425	4C	1
fibroblast growth factor receptor 3 (achondroplasi	FGFR3	2261	NP_075254	NM_000142	4C	
ring finger protein 5	RNF5	6048; 286140	NP_008844	XM_209913	40	
filamin B, beta (actin binding protein 278)	FLNB	2317	NP_001448	NM_001457	40	
eukaryotic translation initiation factor 4A, isofo	EIF4A1	1973	NP_001407	NM_001416	† †	1
TGFB-induced factor 2 (TALE family homeobox)	TGIF2	60436	NP_068581	NM_021809	†	
selective LIM binding factor, rat homolog	SLB	26160	NP_056477	Z012002		1
placental growth factor, vascular endothelial grow	PGF	5228	NP_002623	NIM OTECCS		
ubiquinol-cytochrome c reductase binding protein	UQCRB	7381	NP_006285	NM_003633	40	- 1
high-mobility group box I	HMGB1	3146	NP_002119	NM_00222	t t	fcrh1720
quiescin Q6	QSCN6	200058	MI_00201/		70	fcrh1680
NADH dehydrogenase (ubiquinone) 1 beta subcomplex,	NDUFB4	4/10	NID 007017	NM 003836	4C	fcrb1657
fasciculation and elongation protein zeta 2 (zygin	FEZZ	7710	ND 007075	NM 004547	4C	fcrb1578
Comain containing for transport regulator 6	25.72	0627	NP 005003	NM 005102	4C	fcrb1550
FXYD domain containing in the second of the second	FXYD6	53826	NP_071286	NM_022003	4C	fcrb1547
matrix metalloproteinase 2 (gelatinase A 72). Do	MMP2	4313	NP_004521	NM_004530	4C	fcrb1529
v-erb-b2 erythroblastic leukemia viral oncogena ho	ERBB2	2064	NP_004439	NM_004448	4C	fcrb1528
supervillin	SVIL	6840	NP_068506	NM_021738	4C	fcrb1503

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fcrb2137	4C	NM_000184	NP_000175	3048; 3047	HBG2	h
fcrb2160	4C	NM_003878	NP_003869	8836	GGH	oamma-olutamyi k. d-ol
fcrb2189	4C	NM_015068				Summe-gradiny nydrolase (conjugase, folylpolygamm
fcrb2198	4C	NM_000093	NP_000084	1289;	COL5A1	collagen, type V, alpha I
fcrb2218	4C	NM_005594	NP_005585	4666; 5557	NACA	mascent-nolynopride accomises a contract of the contract of th
fcrb2223	4C	NM_017515	NP_059985	54733	SLC35F2	solute carrier family 25 mounts are applied alpha polyp
fcrb2249	4C	NM_003922	NP_003913	8925	HERC1	heet (homologous to the EZ An Alinea
fcrb2292	4C	NM_012311	NP_036443	22944	KIN	KIN. antigenic determinant of the A
fcrb2305	4C	NM_001402	NP_001393	1915;	EEFIAI	eukaryotic translation elongation factor 1 alpha 1
fcrb2308	4C	NM_005563	NP_981946	3925	STMNI	Stathmin I/oncompation to
fcrb2325	4C	NM_003981	NP_955446		PRC1	Drotein regulator of cytolings; 1
1	4C	NM_024899	NP_079175	79959	C18orf9	chromosome 18 open reading frame o
. [1	NM_002997	NP_002988	6382	SDC1	syndecan 1
10102333	4C	NM_015965	NP_057049	3, 4.	GRIM19	cell death-regulatory protein GRIM19
	4C	NM_017886	NP_060356	54986	FLJ20574	hypothetical protein El 120574
1	4C	NM_007126	NP_009057	7415	VCP	valosin-containing protein
	4C	NM_017777	NP_060247	54903 I	FLJ20345	hypothetical protein FLJ20345
L	4C	NM_080425	NP_536351	2778	GNAS	GNAS complex locus
I	4C	NM_024595	NP_078871	79647 F	FLJ12666	hypothetical protein FI 112666
	4C	NM_006278	NP_006269	6484 S		sialyltransferase 4C (beta-galactoside alpha-23.c
	4C	NM_138425	NP_612434	113246 C	GRCC10	likely ortholog of mouse gene rich cheeser (10 cases)
fcrb2592 4	4C	NM_172020	NP 742017	9883. D		, Some series teneral cluster, C10 ge
forb2506 4				2;	POMIZI	POM121 membrane glycoprotein (rat)
VI_4300010-		NM_003345	NP_919237	7329	UBE2I	ubiquitin-conjugating enzyme E21 (UBC9 homolog, ye
17617906114	2					

ataxin 2 related protein	A2LP	111273	NP_680782	NM_007245	40	CLCCG131
tubulin-specific chaperone d	TBCD	6904	NP_005984	NM_005993	4C	
KIAA1618	KIAA1618	57714	NP_066005	NM_020954	4C	i
WW domain binding protein 2	WBP2	23558	NP_036610	NM_012478	4C	1
glutathione peroxidase 1	GPX1	2876	NP_958799	NM_000581	4C	
protein serine kinase H1	PSKH1	5681	NP_006733	NM_006742	4C	1
ATP-dependent glucokinase	ADP-GK	83440	NP_112574	NM_031284	4C	fcrb3237
hypothetical protein MGC4562	MGC4562	115752	NP_588616	NM_133375	4C	fcrb3227
dermatan 4 sulfotransferase 1	D4ST1	113189	NP_569735	NM_130468	4C	tcrb3169
lipoic acid synthetase	LIAS	11019	NP_919433	NM_006859	4C	fcrb3165
putative dimethyladenosine transferase	HSA9761	27292	NP_055288	NM_014473	4C	tcrb3135
zinc finger, DHHC domain containing 14	ZDHHC14	79683	NP_714968	NM_024630	4C	fcrb3134
synaptonemal complex protein SC65	SC65	10609	NP_006446	NM_006455	40	fcrb3083
serine/arginine repetitive matrix 2	SRRM2	23524	NP_057417	NM_016333	4C	fcrb3073
ring finger protein 149	RNF149	284996	NP_775918	NM_173647	4C	tcrb3017
SH3 domain binding glutamic acid-rich protein like	SH3BGRL3	83442	NP_112576	NM_031286	4C	tcrb2933
collagen, type XII, alpha 1	COL12A1	1303	NP_542376	NM_004370	4C	fcrb2871
profilin I	PFNI	5216	NP_005013	NM_005022	4C	fcrb2813
				NM_018071	4C	fcrb2800
runt-related transcription factor 3	RUNX3	864	NP_004341	NM_004350	4C	fcrb2715
ribosomal protein L28	RPL28	6158	NP_000982	166000 WN	4C	fcrb2713
hemoglobin, alpha 2	HBA2	3040	NP_000508	NM_000517	4C	fcrb2704
sema domain, seven thrombospondin repeats (type 1	SEMA5A	9037	NP_003957	NM_003966	4C	fcrb2690
KIAA0663 gene product	KIAA0663	9877	NP_055642	NM_014827	4C	fcrb2658
neural precursor cell expressed, developmentally d	NEDD5	4735; 55917 NEDD5	NP_004395	NM_004404	4C	fcrb2635

serine protease inhibitor, Kunitz type, 2	SPINT2	10653 S	NP_066925	1404 021102		
ribosomal protein L12	RPL12		NF_00096/		4C	- 1
calcium/calmodulin-dependent protein kinase kinase	Į Ž		NID 000067		4C	fcrb4388 4
CITAL LINV KHIGOV			NP 757380	NM 006549	4C	fcrb4378 2
SNF1-like kinase	SNFILK	150094	NP_775490	NM_173354	4C	fcrb4375
hypothetical protein MGC14327	MGC14327	94107	NP_444273	NM_053045	4C	
golgi autoantigen, golgin subfamily a. 2	GOLGA2	2801	NP_004477	NM_004486	4C	
NADH dehydrogenase (ubiquinone) Fe-S protein 8 23	NDUFS8	4728	NP_002487	NM_002496	+	
ankyrin repeat and SOCS box-containing 3	ASB3	51130	NP_665862	NM_016115	700	ı
pleckstrin homology domain containing, family J me	PLEKHJI	55111; 400670	NP_060519	NM_018049	4C	_
tubulin, alpha 3	TUBA3	7846	NP_006000	NM_006009	4C	
ribosomal protein S20	RPS20	6224	NP_001014	NM_001023	46	1
ubiquitin specific protease 33	USP33	23032	NP_963920	/10C10 MM		1
ribosomal protein S3	RPS3	6188; 6079; 114599	NP_000996	C001002	<u>}</u>	1
chromosome 14 open reading frame 169	C14orf169	79697	NP_078920	NM_024644	†C	fcrh4738
ribosomal protein S8	RPS8	6202; 94161	NP_001003	NM_001012	ĵ t	forb/100
VAMP (vesicle-associated membrane protein)-associa	VAPA	9218	NP_919415	NM_001012	1	forh3841
Finkel-Biskis-Reilly murine sarcoma virus (FBR-MuS	FAU	2197	NP_001988	NM_001997	AC +	fcrh3725
hypoxia inducible factor 3, alpha subunit	HIF3A	64344	NP_690009	NM_152794	S &	forh3718
prothymosin, alpha (gene sequence 28)	PTMA	5757; 51685 PTMA	NP_002814	NM_002823	2 5	fcrh3704
discs, large homolog 7 (Drosophila)	DLG7	9787	NP_055565	NM_00202	AC +C	forh3702
follistatin-like 1	FSTL1	111167	NP_009016	NM_007085	2 4	forb3654
2 DKFZP586H2123 protein	DKFZP586H2 123	25891	NP_056245	NM_015430	1	forb2651
Williams Beuren syndrome chromosome region 20C	WBSCR20C	260294; 155400; 3403	NP_684281	NM_148980	7 6	forh3579 40
					Ŝ	forh 2510

mortality factor 4 like 1	MORF4L1	10933	NP_996670	NM_206839		
butyrophilin, subfamily 3, member A1	BTN3A1	111119	NP_919423	00/00 MIN		
ankyrin repeat and BTB (POZ) domain containing 1	ABTBI	80325	NP_/42025	NM 007048		
snail homolog 2 (Drosophila)	SNAI2	6591	NP_003039	NM 032548		1
protein phosphatase 1, regulatory (inhibitor) subu	PPP1R12C	54776	NP_060077	NM 003068		
ribosomal protein L13	RPL13	6137	NP_150254	NM_033251		forbss36 40
hypothetical gene LOC128439	LOC128439	128439	NP_620585	NM_139016	40	
				NM_018630	4C	
hypothetical protein FLJ20308	FLJ20308	54890	NP_060228	NM_017758	4C	
septin 6	6-Sep	23157	NP_665801	NM_145799	† *	.1
hypothetical protein FLJ36674	FLJ36674	284040	NP_775893	NM_1/3622	†	
nuclear cap binding protein subunit 2, 20kDa	NCBP2	22916	NP_031388	NIM_00/362	5 6	
ribosomal protein L28	RPL28	6158	NP_000982	NIM 007272	30	1
insulin induced gene 2	INSIG2	51141	NP_057217	NIM 000001	40	fcrb5214
G-rich RNA sequence binding factor 1	GRSF1	2926	NP_002083	NIM 016122	40	- 1
MGC4170 protein	MGC41/0	771.70	NID 000000	COCCOO MIN	4C	fcrb5177
i roosoniai protein, iarge, PO	MCCAITO	70158	NP 077288	NM_024312	4C	fcrb5092
ribosomal protein laws po	RPLP0	6175	NP_444505	NM_053275	4C	fcrb4781
proteasome (prosome macronsin) 265	PSMD8	5714	NP_002803	NM_002812	4C	1
cell death-regulatory protein GRIM19	GRIM19	31079; 148113; 37488	NF_03/049			1
ferritin, heavy polypeptide 1	FTHI	2495	NP_002023	NW 015055	4C	fcrb4696
LSM4 homolog, U6 small nuclear RNA associated (S.	LSM4	23804	VIII 000000	NM 000032	4C	fcrb4599
short coiled-coil protein	SCOC	26007	ND 036453	NM 012321	4C	fcrb4570 4C
chromosome 20 open reading frame 104	C20orf104	51230	ND 115026	NM 032547	4C	- 1
hypothetical protein MGC4368	MGC4368	79415	08/8/0_4M	NM 016436	4C	fcrb4428
			VID OTOTO	NM 024510	4C	fcrb44 5

zinc finger protein 36, C3H type-like 1	ZFP36L1	677 Z	NP_004917			L
tetraspanin similar to TM4SF9	DC-TM4F2	19	NP_112189	NM 004026		
ATP synthase, H+ transporting, mitochondrial F0 co	ATP51	521 A	NP_009031	NM_007100		fcrb6896 4C
zinc finger protein 423	ZNF423	23090 2	NP_055884	NM_015069		fcrb6834 4C
copine II	CPNE2	221184	NP_689940	NM_152727		1
sno, strawberry notch homolog 1 (Decombined)	SBNOI	55206	NP_060653	NM_018183		1
EphB3	ЕРНВ3	2049	NP_004434	NM_004443		1
ceroid-linofuscinosis normanila	CLN3	1201	NP_000077	NM_000086	С	icrb6/4/ 4C
WD repeat and SOCS how contributed	WSBI	26118	NP_599027	NM_134264	4C	ı
C-terminal binding protein 1	CTBP1	1487	NP_001319	NM_001328	40	
hypothetical protein FLJ22329	FLJ22329	79709	NP_078932	INM_024656		
potassium voltage-gated channel, shaker-related su	KCNAB3	9196	NP_004723	NIM 024/32	46	
SWI/SNF related, matrix associated, actin dependen	SMARCC2	6601	NF_020/06	NM 00732	4C	fcrb6460
hsp70-interacting protein	HSPBPI	23040	NID (2020)	NM 003075	4C	fcrb6436
Sit 3 dollall containing, Ysc84-like I (S. cerevisiae		22640	NP 036300	NM 012267	4C	fcrb6432
SH3 domain (NNA) II (DNA directed) polypeptide G	SH3YL1	26751	NP_056492	NM_015677	4C	fcrb6251
nolymerase (PNA) if (PNA) is	POLR2G	5436	NP_002687	NM_002696	4C	I
Dolvmerase (DNA dimension on talling 4	POLDIP2	26073	NP_056399	NM_015584	4C	1
zinc finger DHHC domain	ZDHHC4	55146	NP_060576	NM_018106	4C	tcrb6102
general franscription forth IIIC	GTF3C5	9328	NP_036219	NM_012087	4C	Icrb6062
seed family domain containing	SCFD2	152579	NP_689753	NM_152540	40	100012
caspase 8, apoptosis-related cystoine protoco	CASP8	841	NP_203522	NM_033357	7	fark6010
chromosome 20 open reading frame 31	C20orf31	55741	NP_060687	/17010 MIN	40	fcrb6009
PP784 protein	PP784	1114932	NP_982287	NM 018217	4C	fcrb5928
ankyrin repeat domain 10	ANKRD10	35608	NF_060134	CSPEUC WN	4C	fcrb5751
		22/00	ND 060124	NM 017664	4C	fcrb5702
				NM_017710	4C	tcrb5675

to to<							
88 4C NM 032302 NP_115678 84262 MGC10911 hypothetical peroxifedoxin 3 4C NM_053048 NP_653623 121260; SLC15A4 solute carrier 3 4C NM_053048 NP_859428 7001 PRDX2 peroxifedoxin 9 4C NM_016467 51240 LOC51240 hypothetical p 2 4C NM_0174923 NP_06323 54957 DLP Dim1-like pro 3 4C NM_03393 NP_06323 54957 DLP Dim1-like pro 4 4C NM_03393 NP_06323 54957 DLP Dim1-like pro 4 4C NM_03393 NP_06323 54957 DLP Dim1-like pro 4C NM_03394 NP_06323 54957 DLP Dim1-like pro 4C NM_003746 NP_033377 8655 DNCL1 dynein, cytopl 4C NM_004785 NP_035205 3106 HLA-B major histocor 4C N	tropomyosin 3			NP_705935			1
88 4C IMM_032302 INP_115678 84.262 MGC10911 hypothetical protein	tubulin, gamma complex associated protein 3			NP_006313			
88 4C NM_032302 NP_115678 84262 MGC10911 hypothetical peroxiredoxin 36 4C NM_033048 NP_663623 121260; SLC15A4 solute carrier 31 4C NM_005809 NP_859428 7001 PRDX2 peroxiredoxin 4C NM_016467 S1240 LOC51240 hypothetical p 4C NM_0174923 NP_060323 54957 DLP Dim1-like pro 4C NM_00393 NP_003984 1196 CLK2 CDC-like kina 4C NM_00393 NP_0039377 8655 DNCL1 dynein, cytopl 4C NM_003746 NP_003776 9351 SLC9A3R2 solute carrier f 4C NM_004785 NP_003137 6749 SSRP1 structure speci 4C NM_001667 NP_003137 6749 SSRP1 structure speci 4C NM_008629 NP_002759 5119 PCOLN3 procollagen (ty 4C NM_008629 NP_242197 1301	hepatitis C virus core-binding protein 6			NP_076423			
8 4C NM_032302 NP_115678 84262 MGC10911 hypothetical protein processor 6 4C NM_033048 NP_663623 121260; SLC15A4 solute carrier 3 4C NM_03809 NP_859428 7001 PRDX2 peroxiredoxin 1 4C NM_016467 51240 LOC51240 hypothetical processor 2 4C NM_0174923 NP_060323 54957 DLP Dim1-like processor 3 4C NM_003993 NP_060323 203260; MGC31967 hypothetical processor 4 4C NM_003746 NP_003999 5034 P4HB procollagen-processor 8 4C NM_003746 NP_003737 8655 DNCL1 dynein, cytopl 5 4C NM_003146 NP_003137 6749 SSRP1 major histocor 6 4C NM_003146 NP_035205 3106 HLA-B major histocor 6 4C NM_00354 NP_035205 25948 CCT3 chaperonin cor 6 4C NM_001697 NP_001688 539 AT	eukaryotic translation initiation factor 4E-like 3			NP_004837			
8 4C INM_032302 INP_115678 84262 MGC10911 hypothetical provincedoxin 6 4C NM_053048 NP_663623 121260; SLC15A4 solute carrier 3 4C NM_05809 NP_859428 7001 PRDX2 peroxiredoxin 1 4C NM_016467 51240 LOC51240 hypothetical proxiredoxin 2 4C NM_0174923 NP_060323 54957 DLP Dim1-like proxiredoxin 3 4C NM_003993 NP_003323 54957 DLP Dim1-like proxiredoxin 4 4C NM_003993 NP_003984 1196 CLK2 CDC-like kina 4 4C NM_003746 NP_003777 8655 DNCL1 dynein, cytopl 5 4C NM_003746 NP_003137 6749 SSRP1 structure speci 5 4C NM_003746 NP_003137 6749 SSRP1 structure speci 5 4C NM_001268 NP_002759 5119	zinc finger protein 397	ZNF397		NP_115723			
8 +C INM_032302 INP_115678 84262 MGC10911 hypothetical provinced pro	polo-like kinase 1 (Drosophila)	PLKI		NP_005021			1
8 +C NM_032302 NP_115678 84262 MGC10911 hypothetical processor 6 4C NM_033048 NP_663623 121260; SLC15A4 solute carrier 3 4C NM_05809 NP_859428 7001 PRDX2 peroxiredoxin 1 4C NM_016467 51240 LOC51240 hypothetical proxiredoxin 2 4C NM_0174923 NP_060323 54957 DLP Dim1-like proxiredoxin 3 4C NM_0174923 NP_060323 203260; MGC31967 hypothetical proxiredoxin 4 C NM_00393 NP_060323 203260; MGC31967 hypothetical proxiredoxin 4 C NM_00393 NP_003984 1196 CLK2 CDC-like kine 4 C NM_003746 NP_003737 8655 DNCL1 dynein, cytopl 4 C NM_003746 NP_003737 8655 DNCL1 dynein, cytopl 4 C NM_00374 NP_003137 6749 SSRP1 structure speci 4 C NM_003629 NP_036205 22948	origin recognition complex, subunit 5-like (yeast)	ORC5L		NP_859531			ı
8 4C NM_032302 NP_115678 84262 MGC10911 hypothetical processor 6 4C NM_033048 NP_663623 121260; SLC15A4 solute carrier 3 4C NM_005809 NP_859428 7001 PRDX2 peroxiredoxin 4C NM_016467 51240 LOC51240 hypothetical processor 4C NM_0174923 NP_060323 54957 DLP Dim1-like processor 4C NM_003993 NP_003984 1196 CLK2 CDC-like kina 4C NM_003746 NP_003737 8655 DNCL1 dynein, cytopl 4C NM_004785 NP_003737 8655 DNCL1 dynein, cytopl 4C NM_004785 NP_003137 6749 SSRP1 structure specing 4C NM_001697 NP_0036205 22948 CCT5 Chaperonin condition	dimethylarginine dimethylaminohydrolase 2	DDAH2		NP_039268			1
8 4C NM_032302 NP_115678 84262 MGC10911 hypothetical protein live carrier 6 4C NM_033048 NP_663623 121260; SLC15A4 solute carrier 3 4C NM_005809 NP_859428 7001 PRDX2 peroxiredoxin 4C NM_016467 NP_060323 54957 DLP Dim1-like pro 4C NM_174923 NP_777383 203260; MGC31967 hypothetical p 4C NM_003993 NP_003984 1196 CLK2 CDC-like kina 4C NM_003746 NP_003737 8655 DNCL1 dynein, cytopl 4C NM_003746 NP_003737 8655 DNCL1 dynein, cytopl 4C NM_003746 NP_003737 8655 DNCL1 dynein, cytopl 4C NM_003746 NP_003776 9351 SLC9A3R2 solute carrier f 4C NM_003746 NP_003137 6749 SSRP1 structure speci 4C NM_002768 NP_001688	collagen, type XI, alpha 1	COLHAI		NP_542197	NM_080629	#C	1
8 4C NM_032302 NP_115678 84262 MGC10911 hypothetical prothetical processor 6 4C NM_033048 NP_663623 121260; SLC15A4 solute carrier 3 4C NM_005809 NP_859428 7001 PRDX2 peroxiredoxin 1 4C NM_016467 51240 LOC51240 hypothetical proxiredoxin 2 4C NM_0174923 NP_060323 54957 DLP Dim1-like proxiredoxin 3 4C NM_00393 NP_003984 1196 CLK2 CDC-1ike kina 4 C NM_003993 NP_003984 1196 CLK2 CDC-like kina 4 C NM_003746 NP_003737 8655 DNCL1 dynein, cytopl 4 C NM_003746 NP_003736 9351 SLC9A3R2 solute carrier f 4 C NM_003146 NP_003137 6749 SSRP1 structure speci 4 C NM_0012073 NP_036205 22948 CCT5 chaperonin cor	procollagen (type III) N-endopeptidase	PCOLN3		NP_002759	NM_002768	40	1
8 4C NM_032302 NP_115678 84262 MGC10911 hypothetical 5 4C NM_033048 NP_663623 121260; SLC15A4 solute carrier 3 4C NM_005809 NP_859428 7001 PRDX2 peroxiredoxi 4C NM_016467 S1240 LOC51240 hypothetical 4C NM_0174923 NP_060323 54957 DLP Dim1-like pr 4C NM_003993 NP_003984 1196 CLK2 CDC-like kin 4C NM_003918 NP_003737 8655 DNCL1 dynein, cytop 4C NM_003746 NP_003736 9351 SLC9A3R2 solute carrier 4C NM_003146 NP_003137 6749 SSRP1 structure spec 4C NM_012073 NP_036205 22948 CCT5 chaperonin co	ATP synthase, H+ transporting, mitochondrial F1 co	ATP5O	539	NP_001688	NM_001697	† c	1
8 4C NM_032302 NP_115678 84262 MGC10911 hypothetical 5 4C NM_032302 NP_663623 121260; SLC15A4 solute carrier 3 4C NM_005809 NP_859428 7001 PRDX2 peroxiredoxin 4 C NM_016467 NP_060323 54957 DLP Dim1-like proxiredoxin 4 C NM_0174923 NP_777583 203260; MGC31967 hypothetical 4 C NM_003993 NP_003984 1196 CLK2 CDC-like kin 4 C NM_003746 NP_003737 8655 DNCL1 dynein, cytop 4 C NM_005514 NP_005505 3106 HLA-B major histoco 4 C NM_003146 NP_003137 6749 SSRP1 structure spec	chaperonin containing TCP1, subunit 5 (epsilon)	CCT5		NP_036205	NM_012073	6 6	1
4C NM_032302 NP_115678 84262 MGC10911 hypothetical 4C NM_053048 NP_663623 121260; SLC15A4 solute carrier 4C NM_005809 NP_859428 7001 PRDX2 peroxiredoxi 4C NM_016467 NP_060323 54957 DLP DLP Dim1-like proxiredoxi 4C NM_174923 NP_777583 203260; MGC31967 hypothetical 4C NM_003993 NP_003984 1196 CLK2 CDC-like kin 4C NM_003746 NP_003737 8655 DNCL1 dynein, cytop 4C NM_004785 NP_005505 3106 HLA-B major histoco 4C NM_004785 NP_004776 9351 SLC9A3R2 solute carrier	structure specific recognition protein 1	SSRP1	6749	NP_003137	NM_003146		. I
4C NM_032302 NP_115678 84262 MGC10911 5 4C NM_0323048 NP_663623 121260; SLC15A4 8 4C NM_005809 NP_859428 7001 PRDX2 4C NM_016467 51240 LOC51240 4C NM_0174923 NP_060323 54957 DLP 4C NM_003993 NP_003984 1196 CLK2 4C NM_003746 NP_003737 8655 DNCL1 4C NM_005514 NP_005505 3106 HLA-B	solute carrier family 9 (sodium/hydrogen exchanger	SLC9A3R2	9351	NP_004776	NM_004/85	3 8	fork 7820
4C NM_032302 NP_115678 84262 MGC10911 6 4C NM_053048 NP_663623 121260; SLC15A4 4C NM_005809 NP_859428 7001 PRDX2 4C NM_017853 NP_060323 51240 LOC51240 4C NM_174923 NP_777583 203260; MGC31967 4C NM_003993 NP_003984 1196 CLK2 4C NM_003746 NP_003737 8655 DNCL1	S.	HLA-B	3106	NP_005505	NM_005514	40	1010/010
4C NM_032302 NP_115678 84262 MGC10911 4C NM_053048 NP_663623 121260; SLC15A4 4C NM_005809 NP_859428 7001 PRDX2 4C NM_016467 51240 LOC51240 4C NM_017853 NP_060323 54957 DLP 4C NM_174923 NP_777583 203260; MGC31967 4C NM_003993 NP_003984 1196 CLK2 4C NM_000918 NP_000909 5034 P4HB	dynein, cytoplasmic, light polypeptide 1	DNCLI	8655	NP_003737	NM_003746	4C	E-17717
4C NM_032302 NP_115678 84262 MGC10911 hypothetical pothetical pothet	procollagen-proline, 2-oxoglutarate 4-dioxygenase	Р4НВ	5034	NP_000909	NM_000918	4C	tcrb7584
4C NM_032302 NP_115678 84262 MGC10911 hypothetical pothetical pothet	CDC-like kinase 2	CLK2	1196	NP_003984	NM_003993	4C	Icrb /453
4C NM_032302 NP_115678 84262 MGC10911 hypothetical potentical potent	hypothetical protein MGC31967	MGC31967	203260; 84904	NP_777583	NM_174923	†	6 1717
4C NM_032302 NP_115678 84262 MGC10911 hypothetical Intercell 4C NM_053048 NP_663623 121260; 114130 SLC15A4 solute carrier 4C NM_005809 NP_859428 7001 PRDX2 peroxiredoxir 4C NM_016467 51240 LOC51240 hypothetical Intercell	Dim1-like protein	DLP	54957	NP_060323	NM_01/853	10 to	fa-k7202
4C NM_032302 NP_115678 84262 MGC10911 hypothetical peroxiredoxir 4C NM_053048 NP_663623 121260; 114130 SLC15A4 solute carrier 4C NM_005809 NP_859428 7001 PRDX2 peroxiredoxir		LOC51240	51240		NM_016467	1 1 1	forb7320
4C NM_032302 NP_115678 84262 MGC10911 hypothetical NM_053048 NP_663623 121260; SLC15A4 solute carrier	peroxiredoxin 2	PRDX2	7001	NP_859428	NM_005809	40	fa-67221
4C NM_032302 NP_115678 84262 MGC10911		SLC15A4	121260; 114130	NP_663623	NM_053048	7	fa-67112
	hypothetical protein MGC10911	MGC10911	84262	NP_115678	NM_032302	45	forb7026

	CELL	1073	NP 005498	/00000/		
tropomyosin 3	TPM3		NF_/05935	NIM 005507		
GABA(A) receptor-associated protein-like 2	RAPL2	5	NIP_202025	NM 153649		
ubiquitin-conjugating enzyme E2, J2 (UBC6 homolog,		118424	ND 000317	NM 007285		fcrb9639 4C
hypothetical protein FLJ20718	8	35027	NP 010440	NM 194458		
chromosome 9 open reading frame 95		54981	NP_060351	NM 017030		- 1
neighbor of COX4	NOC4	10328	NP_006058	NIM 017881		
prosaposin (variant Gaucher disease and variant me	PSAP	5660	NP_002769	NM 006067	C	
LSM5 homolog, U6 small nuclear RNA associated (S.	LSM5	23658	NP_036454	NM 002728	C	
Dicer1, Dcr-1 homolog (Drosophila)	DICERI	23405	NP_805187	1012222 MN 012222	4C	- 1
HLA-B associated transcript 3	BAT3	7917	NP_542434	NM 177438	4C	- 1
CXXC finger 5	CXXC5	51523	NP_057547	NM_016463	40 +6	
necdin homolog (mouse)	NDN	4692	NP_002478	INM_00248/		
nucleoporin 62kDa	NUP62	23636	NP_714941	NIM 002102	40	
Thy-1 cell surface antigen	5 THYI	7070; 94105 THY1	NP_006279	MIN 152210	40	
candidate mediator of the p53-dependent G2 arrest	REPRIMO	204/2	1NI _002019	OCCCO MIN	4C	fcrb8940
02(KN02) Small nuclear RNA auxiliary factor 1	t t	20173	OI &CYU din	NM 019845	4C	fcrb8908
113/BNI 123 Element binding transcription fa	112AF1	7307	NP_006749	NM_006758	4C	fcrb8901
sterol regulations of the state	SREBF2	6721	NP_004590	NM_004599	4C	tcrb8877
hypothetical protein FI 11/1152	FLJ14153	64747	NP_073573	NM_022736	+	1
HSPC157 protein	HSPC157	29092		NM_0141/9		1
SHC (Src homology 2 domain containing) transformin	SHCI	6464	NP_892113	NIM 014120	40	fcrb8740
interferon regulatory factor 1	IRF1	3659	NF_002189	NM 002020	40	fcrb8668
clusterin (complement lysis inhibitor, SP-40,40, s	CLU	11191	NF_9/6084	801C00 MN	4C	fcrb8664
autism susceptibility candidate 2	AUTS2	20000	ND 07/000	02220C MN	4C	fcrb8516
ilypoinetical protein FLJ10377	* F2 100 / /	36063	ND 056385	NM 015570	4C	fcrb8504
hymoth xi - 1	FL110377	55131	NP_060547	NM_018077	4C	tcrb8465

2)			_		(
10107/14	+	NM_133480	NP_597815	10474	TADA3L	transcriptional adaptor 3 (NGG) hamolog veget) 1:
tcrb9720	4C	NM 000194	NP 000185	13051	HDDTI	i (Coo de monoros, yeast)-II
fcrh9751	20	10000	701000	1070	HPRII	hypoxanthine phosphoribosyltransferase (Lesch-Ny
1077731	40	NM_198335	NP_938149	23193;	GANAB	glucosidase, alpha; neutral AB
fcrb9843	4C	NM 003407	NID 002200	7530		
fcrh9856	40	NIA 015250	NF_003398	/538	ZFP36	zinc finger protein 36, C3H type, homolog (mouse)
fa-b0071	\$ 7	INMI_015350	NP_056165	23507	TA-LRRP	T-cell activation leucine repeat-rich protein
107071	#	NM_032800	NP_116189	84886	FLJ14525	hypothetical protein FLJ14525
rcrb9959	4C	NM_018462	NP_060932	55845	C3orf10	Chromosome 2 case and the C
fcrc0075	4C	NM_000123	NP 000114	2073	ERCCS	organic o oben regarilă itame 10
fcrc0094	4C	NM 016495	ND 057570	6105	n co	excision repair cross-complementing rodent repair
1	40	NIN 022405	111 00/0/7	31230	1BCID7	TBC1 domain family, member 7
1	Ò	NW_033403	NP_208384	85441	PRIC285	peroxisomal proliferator-activated receptor A inte
1	+	NM_020457	NP_065190	57215	THAPII	THAP domain containing 11
Icrc0241	4°C	NM_006612	NP_006603	10749	KIFIC	kinesin family member 1C
fcrc0295	4C	NM_003618	NP_003609	8491	MAP4K3	milogen-activated protein Lincoln
fcrc0350	4C	NM_015180	NP 878918	23224	SVNES	Son were already will ask kinase kinase kin
fere0379	4C	NIM 002607	7 In 2007		0.11402	speculii repeat containing, nuclear envelope 2
		1805007	NP_003678	8572	PDLIM4	PDZ and LIM domain 4
	4C	NM_002809	NP_002800	5709	PSMD3	proteasome (prosome, macronain) 268 submit non A
fcrc0430	4C	NM_002778	NP_002769	5660	PSAP	prosaposin (variant Caucher dieses)
fcrc0487	4C	NM 002231	CCCCOO GN	2720	17 4 1 1	r : composition (variatit Gaucher disease and variant me
	4C	NIA 015275		3/32	KAII	kangai 1 (suppression of tumorigenicity 6, prostat
		C/ 5C10 TAINI	NP_955749	25778	DustyPK	dusty protein kinase
L	4C	NM_005167	NP_786886	389; 333926 RHOC		ras homolog gene family member C
fcrc0559 4	4C	NM_001824	NP_001815	1158	CKM	creatine kinase muscle
fcrc0591 4	4C	NM_001686	NP 001677	506	ATPSR	ATD conthoco III
fcrc0651 4	4C	NM 004518	NP 742107	3785		millocnondrial F1 co
					VCINO?	potassium voltage-gated channel, KQT-like subfamil
NIVI ADDODIO	•					

202333

CMYA5

1213; 29045 CLTC

clathrin, heavy polypeptide (Hc)

cardiomyopathy associated 5

hypoxia up-regulated 1

fcrc0959

fcrc0839

NM_001859 NM_006389 NM_004859 NM_153610

NP_001850

1317 10525

SLC31A1

solute carrier family 31 (copper transporters), me

fa

HYOUI

NP_006380 NP_004850 NP_705838

fcrc0771 |4C fcrc0695 |4C fcrc0654 | 4C

From Figure 2 (MAY-4880Clated)						
nucleolar protein family 6 (RNA-associated)	NOL6	65083	NP_631981	NM_022917	4C	fcrc5671
ribosomal protein L12	RPL12	6136	NP_000967	NM_000976	4C	fcrc5577
				NM_022359	4C	fcrc5516
zinc finger, FYVE domain containing 16	ZFYVE16	9765	NP_055548	NM_014733	4C	L
actin binding LIM protein 1	ABLIMI	3983	NP_006711	NM_002313	40	
DiGeorge syndrome critical region gene 8	DGCR8	54487; 128989	NP_073557	NM_152906	4C	
hypothetical protein MGC5508	MGC5508	79073	NP_076997	NM_024092	4C	1
splicing factor, arginine/serine-rich 12	SFRS12	140890	NP_631907	NM_139168	4C	1crc4896
phosphoinositide-3-kinase, regulatory subunit, pol	PIK3R2	5296	NP_005018	NM_005027	4C	fcrc4658
integrin, alpha 6	ITGA6	3655	NP_000201	NM_000210	40	Icrc4485
ubiquitin-conjugating enzyme E2G 1 (UBC7 homolog,	UBE2G1	7326	NP_872623	NM_003342	4C	fcrc4390
APG16 autophagy 16-like (S. cerevisiae)	APG16L	55054	NP_942593	NM_030803	4°C	Icrc4360
WD40 protein Ciao l	CIA01	9391	NP_004795	NM_004804	4C	tcrc4054
centaurin, gamma 3	CENTG3	116988	NP_114152	NM_031946	4C	tcrc4045
hypothetical protein FLJ20255	FLJ20255	54868	NP_060198	NM_017728	4C	tcrc3993
DEAH (Asp-Glu-Ala-His) box polypeptide 15	DHX15	1665	NP_001349	NM_001358	4C	tere3750
galactosamine (N-acetyl)-6-sulfate sulfatase (Morq	GALNS	2588	NP_000503	NM_000512	4C	fcrc3739
				NM_144983	4C	fcrc3415
NADH dehydrogenase (ubiquinone) 1, subcomplex unkn	NDUFC2	4718	NP_004540	NM_004549	4C	fcrc3358
polymerase I and transcript release factor	PTRF	284119	NP_036364	NM_012232	4C	fere3240
zinc finger protein 565	ZNF565	147929	NP_689690	NM_152477	4C	icrc3048
arginyl aminopeptidase (aminopeptidase B)	RNPEP	6051	NP_064601	NM_020216	6	rere3009
zinc finger protein 62 homolog (mouse)	ZFP62	92379	NP_689496	NM_152283	4C	1crc2997
mitochondrial translation optimization 1 homolog (MTOI	25821	NP_598400	NM_133645	4C	fcrc2969
nucleolar protein family A, member 2 (H/ACA small	NOLA2	55651	NP_060308	NM_017838	4C	fcrc2948
						,

					V	NYI-4308219v1
ras homolog gene family, member E	ARHE	390	NP_005159	NM_005168	4C	htcr0415
transmembrane 4 superfamily member [TM4SF1	4071	NP_055035	NM_014220	†	
serologically defined breast cancer antigen 84	SDBCAG84	51614	NP_938408	NM_015966	4C	
basigin (OK blood group)	BSG	682	NP_940993	NM_001728	45	1
ATP synthase, H+ transporting, mitochondrial F1 co	ATP5A1	498	NP_004037	NM_004046	4C	I
KIAA0922 protein	KIAA0922	23240	NP_056011	NM_015196	46	1
isocitrate dehydrogenase 1 (NADP+), soluble	IDHI	3417	NP_005887	NM_005896	16	
zinc finger protein 593	ZNF593	51042	NP_056955	NM_015871	40	1
suppression of tumorigenicity 13 (colon carcinoma)	ST13	6767	NP_003923	NM_003932	4°C	
chromosome condensation-related SMC-associated pro	CNAPI	9918	NP_055680	NM_014865	4C	1
retinoblastoma binding protein 6	RBBP6	5930	NP_116015	NM_006910	4C	1
epsin 2	EPN2	22905	NP_683723	NM_014964	4°C	Icrc /04 /
F-box only protein 7	FBXO7	25793	NP_036311	NM_012179	4C	1crc6990
paired basic amino acid cleaving system 4	PACE4	5046	NP_612198	NM_002570	÷ 6	1crc69/0
chromosome 1 open reading frame 33	C1orf33	51154	NP_057267	NM_016183	#	f (070
neogenin homolog 1 (chicken)	NEOI	4756	NP_002490	NM_002499	\$ 6	fam.6000
egl nine homolog 3 (C. elegans)	EGLN3	112399	NP_071356	NM_022073	100	fcrc6826
LYST-interacting protein LIP8	LIP8	116840	NP_444279	NM_053051	40	fcrc6642
similar to rat tricarboxylate carrier-like protein	BA108L7.2	81855	NP_112233	NM_030971	4C	fcrc6609
tousled-like kinase 2	TLK2	11011	NP_006843	NM_006852	4C	tcrc6566
copine I	7 CPNE1	8904; 10137	NP_690908	NM_006047	40	icrebbb5
plexin A2	PLXNA2	5362; 148697	NP_079455	NM_153014		10100000
TH1-like (Drosophila)	THIL	51497	NP_945327	NM_016397	46	for 6225
replication factor C (activator 1) 2, 40kDa	RFC2	5982	NP_852136	NM_181471	1	Crc5846
surfeit 4	SURF4	6836; 401567	NP_149351	NM_033161		fcrc5797

NYI-4308219v1

				NM 033408	4C	htcr1709
Ksp37 protein	KSP37	83888 k	NP_114156	NM_031950	+	
filamin B, beta (actin binding protein 278)	FLNB	2317 F	NP_001448	MM_00145/		i
F-box only protein 34	FBXO34	55030 F		NM_017943	4C	hfcr1641
				XM_065820	4C	
tenomodulin protein	TNMD	64102	NP_071427	NM_022144	4C	
estrogen-related receptor gamma	ESRRG	2104	NP_996318	NM_001438	4C	
Similar to actinin, alpha 4 (LOC344978), mRNA	344978			XM_293669	40	
glycine amidinotransferase (L-arginine:glycine ami	GATM	2628	NP_001473	NM_001482	4	
thyroid hormone receptor interactor 6	TRIP6	7205; 56996 TRIP6	NP_003293	NM_003302	Ď t	1
natriuretic peptide precursor B	NPPB	4879	NP_002512	NM_002521	t t	
eukaryotic translation elongation factor beta 2	EEF1B2	1933	NP_066944	NM_00223	70 7	. 1
transmembrane 4 superfamily member 13	TM4SF13	27075	NP_055214	NM_014399	à t	- 1
annexin A6	ANXA6	309	NP_004024	NM 011222		hfcr1163
interferon induced transmembrane protein 3 (1-8U)	IFITM3	10410	NP_066362	NIM_021034	40	hfcr1137
peroxisome proliferative activated receptor, delta	PPARD	5467	NP_803184	NIM 000238	40	hfcrl 115
ATPase, H+ transporting, lysosomal 21kDa, V0 subun	ATP6V0B		NP_004038	NIM 00404/	40	hfcr1101
karyopherin alpha 2 (RAG cohort I, importin alpha	KPNA2	3838	NP_002257	NM_002266	40	hfcr0829
nuclear protein double minute 1	MDMI	56890	NP_064513	NM_017440	6 6	hfa-0675
protocadherin 10	PCDH10	57575	NP_116586	NM_032961		HC10024
collagen, type VIII, alpha 2	COL8A2	1296	NP_005193	NM_005202		1200121
CGI-12 protein	CGI-12	51001	NP_057026	746C10 MM	20 6	hforfissi
hypothetical protein FLJ20729	FLJ20729	54680	NP_060423	NIM 015043	4 7	hfcr0496
MAX protein	MAX	4149	NP_932061	NIM_145113	40 7	hfcr0489
keratin 10 (epidermolytic hyperkeratosis; keratosi	KRI IO	3038	711-000-112	NIM 145112	40	hfcr0478

				210000		I
KIAA1055 protein	KIAA1055	23102	INF_055694	XM 216866	C	
high-mobility group nucleosome binding domain 1	HMGNI	3150	NP_004936	NM 015079	C	ı
splicing factor, arginine/serine-rich 9	SFRS9		NP_003760	NM 004065		- 1
actin, beta	ACTB	60; 71	NP_001092	MM 002701		
insulin-like growth factor binding protein 4	IGFBP4	3487	NP_001543	NM 001101	40 +6	
phospholipase D3	PLD3	23646	NP_036400	NM_012268	70 45	hfcr2820 4
hypothetical protein MGC4677	MGC4677	112597	NP_443103	NM_052871	Å to	
DNA segment on chromosome X (unique) 9928 expresse	DXS9928E	9130	NP_004690	NW_004699		
laminin receptor 1 (ribosomal protein SA, 67kDa)	LAMRI	3921	NP_002286	NM_002295	40 +6	1
protein disulfide isomerase related protein (calci	ERP70	9601	NP_004902	NM_004911	à t	. 1
a disintegrin and metalloproteinase domain 12 (me)	ADAM12	8038	NP_067673	NM_003474	Ď ŧ	hfcr2544
heterogeneous nuclear ribonucleoprotein M	HNRPM	4670	NP_112480	NM_005968	4C	
				NM_014054	4C	.1
synaptonemal complex protein SC65	SC65	10609	NP_006446	NM_006455	4C	
early B-cell factor	EBF	1879	NP_076870	NM_024007	4C	1
- A TOOMACKOPIOKEIII AZ/BI				NM_014860	4C	hfcr2148
	HNRPA2B1	3181	NP_112533	NM_031243	4C	htcr1914
THAP domain containing 7	THAP7	80764	NP_085050	NM_030573	40	1
microtubule-associated protein 4	MAP4	4134	NP_112147	NM_002375	à t	hf2-10/2
translocase of outer mitochondrial membrane 7 homo	TOMM7	54543	NP_061932	NM_019059	t t	hfcr1856
HBxAg transactivated protein 2	XTP2	23215	NP_055987	NM_015172	à t	hfcr1705
adiponectin receptor I	ADIPORI	51094	NP_057083	NM_015999	3 6	hfcr1760
regulator of G-protein signalling 3	RGS3	5998	NP_652760	NM_144488	4C	hfcr1733

G G					-	NYI-4308219v1
likely ortholog of mouse signaling intermediate in	SITPEC	51295	NP_057665	NM_016581		1000000
pyrroline-5-carboxylate reductase 1	PYCRI	5831	NP_/22546	14M 000507	، ار	- 1
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex	NDUFA3	4696	NP_004533	NM_004542		
sushi-repeat-containing protein, X-linked	SRPX	8406	NP_006298	NM_006307	1 0 1 0 1 0	hfcr5181 4
ribosomal protein L14	RPL14	9045	NP_003964	NM_003973	†	
cell division cycle 42 (GTP binding protein, 25kDa	CDC42	998	NP_426359	NM_0444/2	5 6	
integrin, alpha 5 (fibronectin receptor, alpha pol	ITGA5	3678	NP_002196	NM 044422	40	- 1
MCM3 minichromosome maintenance deficient 3 (S. ce	MCM3	4172	NP_002379	NM_002388	40	hfcr4488 2
adenosylmethionine decarboxylase I	AMD1	262	NP_001625	NM_001634	† ÷	1
MLL septin-like fusion	MSF	10801	NP_006631	NM_006640	4C	L
tubulin, beta, 4	TUBB4	10381	NP_006077	NM_006086	#	i
TRK-fused gene	TFG	10342	NP_006061	NM_006070	100	1
jun D proto-oncogene	DUND	3727	NP_005345	NM_005354	45	1
protein kinase C and casein kinase substrate in ne	PACSIN3	29763	NP_057307	NM_016223	45	1
aminopeptidase puromycin sensitive	NPEPPS	9520	NP_006301	NM_006310	7	
poly(rC) binding protein 2	PCBP2	5094	NP_114366	OLOCOO_IAIN	70 6	
fascin homolog 1, actin-bundling protein (Strongyl	FSCN1	6624	NP_003079	NIM 005017	40	
Kruppel-like factor 2 (lung)	KLF2	10365	NP_057354	NIM 002088	40	- 1
leucine rich repeat neuronal 1	LRRNI	57633	NP_065924	NM_020873	4C	hfcr3413
zinc finger, DHHC domain containing 3	ZDHHC3	51304	NP_057682	NM_016598	40	hfcr3/10/
lectin, galactoside-binding, soluble, 1 (galectin	LGALSI	3956; 57026; 23616	NP_002296	1002903		hfaransc
mitochondrial solute carrier protein	MSCP	51312	NP_061049	NM 003305	40 7	hfcr3180
RNA binding protein S1, serine-rich domain	RNPSI	10921	NP_542161	NM_006711	100	hfcr3140
DKFZP434N1 hypothetical protein DKFZp434N1923	DKFZP434; 923	81858	NP_112236	NM_030974	4C	htcr3022

nicro2228	4°C	NM_017991	NP_060461	55683	FLJ10081	hypothetical protein El 110001
hfcr5237	4C	NM_021738	NP_068506	6840	SVIL	supervillin
htcr5383	4C	NM_002398	NP_002389	4211	MEISI	Meis I, myeloid ecotronic viral integration site 1
hfcr5473	40	NM_021128	NP_066951	5441	POLR2L	polymerase (RNA) II (DNA directed) polymentide I
hfcr5601	AC C	NM_002789	NP_002780	5685	PSMA4	proteasome (prosome, macropain) subunit, alpha tvn
hfcr5695	40	NIM 144584	NP_004688	9128	PRPF4	PRP4 pre-mRNA processing factor 4 homolog (yeast)
	2	MM_144584	NP_653185	113802	FLJ30525	hypothetical protein FLJ30525
1	4C	NM_003286	NP_003277	7150	TOPI	topoisomerase (DNA) I
1	4C	NM_145914	NP_666019	7589	ZNF38	zinc finger protein 38
1	4C	NM_001005	NP_000996	6188; 6079;	RPS3	ribosomal protein S3
hfcr6164	4C	NM_003505	NP_003496	8321	FZDI	frizzled homolog 1 (Droscopkile)
	4C	NM_144498	NP_653081	9885	OSBPL2	oxysterol binding protein-like 2
	#	NM_005493	NP_005484	10048	RANBP9	RAN binding protein 9
	40	NM_019105	NP_115859	7148	TNXB	
1	4C	NM_020404	NP_065137	57124	CD164L1	CD164 sialomucin-like 1
1	4C	NM_003977	NP_003968	9049	AIP	aryl hydrocarbon receptor interacting protein
hfcr6500	Ď ŧ	NM_014718	NP_055533	9746	CLSTN3	calsyntenin 3
i	70 70	NM_003397		5420	PODXL	podocalyxin-like
	70 6	NM_020830		57590	WDFYI	WD repeat and FYVE domain containing 1
	7	MM_005809	NP_859428	7001	PRDX2	peroxiredoxin 2
	46	NM_006058	NP_006049	10318	TNIPI	TNFAIP3 interacting protein
1	4C	NM_001997	NP_001988	2197 I	FAU	Finkel-Biskis-Reilly murine sarcoma virus (EDD N. C.
1	4C	NM_012394	NP_036526	5202 I	PFDN2 r	prefoldin 2
	4C	NM_000517	NP_000508	3040 I	HBA2 h	hemoglobin, alpha 2
mcro931 4C		NM_032548	NP_742025	80325 A	ABTBI a	ankyrin repeat and BTB (POZ) domain containing 1
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	PSMB5 pı	5693 P	NP_002788	NM_002797	C	mioa0826 4C
ATP synthase, H+ transporting, mitochondrial F0 co	ATP5GI A	516	NP_005166	NM_005175		m10a0707 4C
				NM_021122	†	
steroid sulfatase (microsomal), arylsulfatase C, i	STS st	412	NP_000342	NW 021122		mioa0647 40
zinc finger protein 626	ZNF626 z	199777	NF_000340	NIM DODOSI	4C	mioa0595 4C
decay accelerating factor for complement (CD55, Cr			NP_000363	NM 145207	4C	mioa0582 4C
putative membrane protein	LOC54499 p	9	NP_061899	NM 002020	4C	mioa0577 4C
polypyrimidine tract binding protein 2	РТВР2		NP_067013	950010 MIN		mioa0535 4C
LIM protein (similar to rat protein kinase C-bindi	LIM	10611	NP_006448	MIN 031100	40	mioa0494 4C
chromosome 9 open reading frame 78		51759; 1861 C9orf78	NP_057604	NM_016482	4C +C	mioa0407 4C
HSPC163 protein	HSPC163	29097	NP_054903	NM_014184	70 30	mina0380 40
protein phosphatase 1, regulatory (inhibitor) subu	PPP1R3C	5507	NP_005389	NM_005398	40 +t	mioa0328 4C
growth hormone inducible transmembrane protein	GHITM	27069	NP_055209	NM_014394	à đ	mica0311
ribosomal protein S24	RPS24	399785	NF_140962		Ô	mina0308 40
cysteine-rich, angiogenic inducer, 61		J+71	Nin 140000	MM 033000	4C	mioa0246 4C
Pase activating prote		3/01	NP 001545	NM 001554	4C	mioa0204 4C
RAS n21 profein activate (CTn	RASAI	5921	NP_072179	NM_002890	4C	mioa0152 4C
coiled-coil-helix-coiled-coil-helix domain instead	CHCHD2	51142	NP_057223	NM_016139	4C	mioa0132 4C
protein tyrosine phosphatase-like (proling instance)	PTPLB	201562	NP_940684	NM_198402	4C	m10a0104 4C
phosphatidylinosítol glycan, class O	PIGQ	1606	NP_683721	NM_004204	100	
nascent-polypeptide-associated complex alpha polyp	NACA	4666; 5557	NP_005585	NM_000946	3 8	hfcr0613
RAD9 homolog A (S. pombe)	RAD9A	5883	NP_004575	NM_00222		hfcr9549
glucosidase, alpha; neutral AB	GANAB	23193; 80789	NP_958149	NIM 004504		hfcr9290
UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase,	B3GALT3	8/06	ND 020110	NM 108225	4C	hfcr8902
chromosome 10 open reading frame 48	C100rr48	2000/0	NID 140250	NM 003781	4C	hfcr7968
carbonic annydrase IX	CIOLEGO	783078	NP 775847	NM 173576	4C	hfcr7966
Corbonic orbital	CA9	768	NP_001207	NM_001216	4 _C	hfcr6932

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chromosome 10 open reading frame 32	C10orf32	119032	NP_653192	144591	(
protein tyrosine phosphatase, receptor type, M	PTPRM	5797; 80041 PTPRM	NP_002836	NIM 144501	40	mioa 1531 4C
CD2-associated protein	CD2AP	23607	NP_036252	NIM 002875	4C	mioa1520/4C
5-methyltetrahydrofolate-homocysteine methyltransf	MTRR	4552	NP_0/6915	010120 MIN	40	mioa1494 4C
methylmalonyl Coenzyme A mutase	MUT	4594	NP_000246	010700 MINI	40	mioa1473 4C
pumilio homolog 2 (Drosophila)	PUM2	23309	NI 000012	NIM DODSS	4C	mioa1445 4C
serine (or cysteine) proteinase inhibitor, clade E	DERVINE	22260	ND 056133	NM 015317	4C	mioa1427 4C
chromosome 9 open reading frame 83	COOLOS	5057	NP 000593	NM 000602	4C	mioal 380 4C
associated (1 NFRSF16) associated	Corres	01515	NP 057494	NM_016410	4C	mioal354 4C
nerve growth factor recentor (TNIED cold)	NGFRAPI	27018	NP_996800	NM_014380	4C	mioal 337 4C
death associated protein 3	DAP3	7818	NP_387506	NM_033657	145	
		317762			Š	min 133
cyclin K	CCNK	64758:	NF_005849			***************************************
core 1 UDP-galactose:N-acetylgalactosamine-alpha-R	CIGALT2	29071	NF_089905	NM 174070	4C	mioa1285 4C
A I Pase, Ca++ transporting, type 2C, member 1	AIF2CI	1000	NID COOCC	C09C51 WN	40	mioa1279 4C
	A TBOCI	27022	NP 055197	NM 014382	5 4C	mioa1276 4C
splicing factor argining/sering rich ?	SFRS2IP	9169	NP_004710	NM_004719	3 4C	mioal 193 4C
collagen, type I, alpha 2	COL I A2	1278	NP_000080	NM_000089	7/4C	m10a1097/4C
syntaxin binding protein 6 (amisyn)	STXBP6	29091	NP_054897	NM_014178	† † †	
chronic myelogenous leukemia tumor antigen 66	CML66	84955	NP_116258	MINI_022099	0 7	mios 1070 AC
proteasome (prosome, macropain) 26S subunit, ATPas	PSMC2	5701	NP_002794	NIM 032860	140	mioa 1071 4C
HMG-box transcription factor 1	HBP1	26959	NP_036389	NM 002002	2 40	mioa1062 4C
sec1 family domain containing 1	SCFD1	23256	NP_6/8233	NM 010257	8 4C	mioa1058 4C
mitochondrial ribosomal protein L42	IVIRPL42	10071	ND 070255	NM 016106	5 4C	mioa1055 4C
Troponial protell L39	MDDIAS	28977	NP 751918	NM 172178	5 4C	mioa1015 4C
ribosomal protein 120	RPL39	6170	NP_000991	0001000 MN)9 4C	mioa0909 4C
glucocorticoid recentor DNA Line:	GRLF1	2909	NP_077318	NM_024342	00 4C	mioa0890 4C
fatty acid hinding protein A adiment	FABP4	2167	NP_001433	NM_001442	34.40	11110a0862 4C

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influenza virus NSIA hinding protein	IVNSIABP i	10625	NP_006460	NIVI_000469		t
dihydropyrimidinase-like 2	DPYSL2	1808	NP_001377			mioa2652 4C
copine III	CPNE3				#C	mioa2580 4C
uisabled flomolog 2, mitogen-responsive phosphoprot			NP 003900	NM 003909	¢C	mioa2537 4C
disabled homel.		1601	NP_001334	NM_001343	4C	m10a2536 4C
nudix (nucleoside dinhosphoto ii.i.	NUDT5	11164	NP_054861	NM_014142	†	7767801111
signal peptidase complex (18kD)	SPC18	23478; 157708	NP_055115	WW 014300	5	mi092522 4C
chronic myelogenous leukemia tumor antigen 66	CML66	84955	NP_116258	NIM 014200	4C	mioa2478 4C
hypothetical protein FLJ90834	FLJ90834	283135	NF_//3849	098CEU WN	4C	mioa2475 4C
by Creatil		54833	ND 7750 10	NM 173578	4C	mioa2421 4C
CGI-30 protein	CGI-30	51611;	NP_057042	NM_015958	4C	m10a2413 4C
recombining hinding protein comparation constraints	RBPSUH	3516	NP_976029	NM_015874	#	· Second
C-type (calcium dependent, carbohydrate-recognition	CLECSFI	10143	NP_005743	7C/C00_MIN	A) 6	mina2333 40
chromosome 13 open reading frame 12	C13orf12	51371	INF_05/016	NIA 005752	40	mioa2327 4C
malate dehydrogenase 1, NAD (soluble)	IHDM	4190	NID 057010	NM 015932	4C	mioa2319 4C
buly(rC) binding protein 2	1 (01.6)		ND DOSONO	NM 005917	34C	mioa2268 4C
2017 Contract Protein 728	PCRP2	5094	NP 114366	NM_005016	34C	mioa2173 4C
Zinc finger protein 360	ZNF258	9204	NP_660353	NM_007167	84C	mioa2158 4C
ribosomal protein Sc	RPS6	6194	NP_001001	NM_001010	6 4C	m10a2156 4C
destrin factin denotypes:	DSTN	11034	NP_006861	NM_006870	2.4C	m10a2072 4C
trafficking protein particle complete	TRAPPC4	51399	NP_057230	NM_016146	340	IIII0a2013 4C
ubiquitin-conjugating enzyme E2 A /B A T/	UBE2A	7319	NP_861442	NM_003336	1 1 1	11.00117
chemokine (C-X-C motif) recentor 4	CXCR4	7852	NP_003458	NM_003467	40	mina1071 40
cullin 3	CUL3	8452	NP_003581	NIN_003165	3 40	mioa 1953 40
Niemann-Pick disease, type C2	NPC2	10577	NP_006423	NIM 003500	240	mioa 1 662 4C
glyoxalase I	GL01	2739	NF_006699	NM 006432	30 4C	mioa1660 4C
solute carrier family 16 (monocarboxylic acid tran	SLC16A1	0000	710 COC 211.	NM 006708	26 4C	mioa1626 4C
		7759	NP 003042	NM 003051	21 4C	mioa1621 4C
				NM_014671	85 4C	mioa1585 4C

chromosome 15 hypothetical ATG/GTP binding protein	LOC56851 cl	56851	NP_064539	NM_020154	†	0a+070 +C
polymerase (RNA) II (DNA directed) polypeptide C.	POLR2C p	5432	NP_116558	NM_032940		mica4004 4C
signal recognition particle receptor, B subunit	SRPRB si	58477	NP_067026	NM_021203		mica4014 4C
protein kinase, cAMP-dependent, regulatory, type I	PRKARIA p	5573	NP_997637	NM_002734		miss4009/4C
RAB32, member RAS oncogene family	RAB32 R	10981	NP_006825	NM_006834		mina/000 4C
phospholipase A2, group IIA (platelets, synovial f	PLA2G2A p	5320	NP_000291	NM_000300		mica 2062 40
solute carrier family 4 (anion exchanger), member	SLC4A1AP s	22950	NP_060628	NW_018158	Ď ŧ	mica 2044 40
cartilage intermediate layer protein, nucleotide p	CILP	8483	NP_003604	NM_003613	5 6	mica3030 4C
yippee protein	CGI-127	51646	NP_057145	NM_016061	40	min 2022 4C
chromosome 3p21.1 gene sequence	HUMAGCGB (29890	NP_037418	NM_013286	4C	m10a5812 4C
spectrin, beta, non-erythrocytic 1		6711; 56969 SPTBNI	NP_842565	NM_003128	40	m10a3 /99 4C
lysosomal-associated protein transmembrane 4 alpha	LAPTM4A	9741	NP_055528	NM_014713	40	Imioas /86/4C
ATP synthase, H+ transporting, mitochondrial F1 co	АТР5В	506	NP_001677	NM_001686	÷ ;	1111043040 40
S100 calcium binding protein A6 (calcyclin)	S100A6	6277; 400825	NP_055439	11 0 1 40Z4		minoscat
syndecan binding protein (syntenin)	SDCBP	6386	NP_005616	MM 014624	40	mioa3629 4C
protein kinase, interferon-inducible double strand		00.0	Nin Oostoo.	NM 005625	4C	mioa3620 4C
"" our 14 open reading frame 129		8575	NP 003681	NM 003690	4C	mioa3514 4C
	C14orf129	51527	NP_057556	NM_016472	4C	mioa3471 4C
PTD012 protein	PTD012	28970	NP_054758	NM_014039	4C	mioa3439 4C
signal recognition particle 19kDa	SRP19	6728; 7310	NP_003126	NM_003135	4C	m10a3395 4C
metastasis suppressor	MTSS1	9788	NP_055566	NM_014751	40	
protein associated with PRK1	AWPI	54469	NP_061879	NM_019006	1 6	micassa
MAD, mothers against decapentaplegic homolog 5 (Dr	MADH5	4090	NP_005894	NM_005903	3 7	mioa3000 +C
nuclear ubiquitous casein kinase and cyclin-depend	NUCKS	64710	NP_073568	NM_022731	240	mioa2001 4C
vacuolar protein sorting 24 (yeast)	VPS24	51652	NP_057163	NM_016079	8 4 C	m10a2818 4C
forkhead box C2 (MFH-1, mesenchyme forkhead 1)	FOXC2	2303	NP_005242	NM_005251	#4C	m10a2 / 88 4C

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AMP reconcive element hinding motion	CREB1	1385	NP 604391	NM_134442	4C	mioa7323 4C
growth hormone receptor	GHR g	2690	NP_000154	NM_000163	4C	mioa7317 4C
U5 snRNP-specific protein, 116 kD	U5-116KD L	9343	NP_004238	NM_004247	40	mioa/299/4C
Rac/Cdc42 guanine nucleotide exchange factor (GEF)	ARHGEF6 F	9459	NP_004831	NM_004840	4C	mioa7140 4C
ribosomal protein S23	RPS23 r	6228	NP_001016	NM_001025	4C	m10a7015 4C
hypothetical protein LOC114971	LOC114971	114971; 4722		NM_004551	4C	m10a6913 4C
hypothetical protein BC016005	LOC129642	129642	NP_620154	NM_138799	4C	m10a6854 4C
ubiquitin carboxyl-terminal hydrolase L5	UCHL5 1	51377	NP_057068	NM_015984	4C	m10a6739 4C
v-fos FBJ murine osteosarcoma viral oncogene homol	FOS	2353	NP_005243	NM_005252	4C	m10a6/38/4C
WD repeat and SOCS box-containing	WSB1	26118	NP_599027	NM_134264	4C	mioa6731 4C
myosin, heavy polypeptide 9, non-muscle	МҮН9	4627	NP_002464	NM_002473	4C	m10a6721 4C
RAB18, member RAS oncogene family	RAB18	22931	NP_067075	NM_021252	4C	m10a6621 4C
heterogeneous nuclear ribonucleoprotein H1 (H)	HNRPH1	3187	NP_005511	NM_005520	4C	m10a6582 4C
histamine N-methyltransferase	HNMT	3176	NP_008826	NM_006895	4C	m10a6476 4C
toll-like receptor 2	TLR2	7097	NP_003255	NM_003264	4C	m10a6418 4C
mitogen-activated protein kinase kinase kinase 2	MAP3K2	10746	NP_006600	NM_006609	4C	mioa6268 4C
nebulette	NEBL	10529	NP_998734	NM_006393	4C	mioa6174 4C
hypothetical protein MGC40053	MGC40053	158947	NP_689796	NM_152583	4C	m10a6130 4C
phosphorylase, glycogen; liver (Hers disease, glyc	PYGL	5836	NP_002854	NM_002863	4C	mioa6093 4C
squalene epoxidase	SQLE	6713	NP_003120	NM_003129	4C	mioa6091 4C
hypothetical protein DKFZp434B227	DKFZp434B2 27	84223	NP_115639	NM_032263	4C	mioa6064 4C
clock homolog (mouse)	CLOCK	9575	NP_004889	NM_004898	4C	m10a6035 4C
zinc finger protein ANC_2H01	ANC_2H01	51193	NP_057415	NM_016331	2/4C	mioa5902 4C
tumor necrosis factor, alpha-induced protein 6	TNFAIP6	7130	NP_009046	NM_007115	5 4C	mioa5836 4C
cyclin-dependent kinase 7 (MO15 homolog, Xenopus l	CDK7	1022	NP_001790	NM_001799	3/4C	mioa5773 4C

NYI-4308219v1

minox7602_4C NM_178191 NP_835498 93974 ATPIFI ATPase inhibitory factor i							
4C NM_178191 NP_835498 93974 ATPIFI ATPase inhit 4C NM_018434 NP_069904 55819 RNF130 ring finger pt 4C NM_004981 NP_069094 55819 RNF130 ring finger pt 4C NM_004981 NP_065090 57092 PCNP Epstein-Barr 4C NM_006304 NP_997386 401388; FLJ42280 FLJ42280 pt C NM_006304 NP_997386 401388; FLJ42280 pt FLJ42280 pt C NM_006304 NP_997386 401388; FLJ42280 pt FLJ42280 pt C NM_006304 NP_997386 401388; FLJ42280 pt FLJ42280 pt C NM_0138931 NP_903884 4388 BCL6 B-cell CLL/ly C NM_0138931 NP_9038854 6388 BCL6 B-cell CLL/ly C NM_004673 NP_908805 79577 HRPT2 hyperparathyr C NM_001421 NP_9014605 71 ACTG1 actin, gamma C <td>nuclear autoantigenic sperm protein (histone-bindi</td> <td>NASP</td> <td></td> <td></td> <td></td> <td>†</td> <td>0,000</td>	nuclear autoantigenic sperm protein (histone-bindi	NASP				†	0,000
IC NM_178191 NP_835498 93974 ATPIEI ATPase inhit IC NM_018434 NP_06904 558.9 RNF130 ring finger pt IC NM_004951 NP_06904 558.9 RNF130 ring finger pt IC NM_004951 NP_065090 57092 PCNP E2F6 E2F transcrip IC NM_006304 NP_997386 401388: 79978; FLJ42280 pt FLJ42280 pt IC NM_006304 NP_997386 401388: 79978; FLJ42280 pt FLJ42280 pt IC NM_006304 NP_997386 401388: 79978; FLJ42280 pt FLJ42280 pt IC NM_006923 NP_008854 6388 SDF2 stromal cell-d IC NM_006923 NP_078805 79577 HRPT2 hyperparathyr IC NM_006923 NP_001605 71 ACTG1 actin, gamma IC NM_0021431 NP_001605 71 ACTG1 actin, gamma IC NM_002709 NP_001422 2037 EPB	DKFZP564G2022 protein	DKFZP564G2 022					100000
IC NM_178191 NP_835498 93974 ATPHFI ATPase inhit IC NM_018434 NP_060904 55819 RNF130 ring finger proposed in the protein plant of the pro	ubiquitin-like 5	UBL5	59286; 5300	NP_077268			01000000
IC NM_178191 NP_835498 93974 ATPIEI ATPase inhil C NM_018434 NP_060904 55819 RNF130 ring finger pn C NM_004951 NP_060904 55819 RNF130 ring finger pn C NM_004951 NP_065090 1876 E2F6 E2F transcrip C NM_006304 NP_997386 401388; FLJ42280 FLJ42280 pr C NM_006304 NP_997386 401388; FLJ42280 FLJ42280 pr C NM_138931 NP_620309 604; 389185 BCL6 B-cell CLL/ly C NM_006923 NP_008854 6388 SDF2 stromal cell-d C NM_006923 NP_078805 79577 HRPT2 hyperparathyr D NM_004529 NP_001605 71 ACTG1 actin, gamma NM_001614 NP_001605 71 ACTG1 actin, gamma NM_007271 NP_001605 71 ACTG1 actin, gamma NM_001421 NP_0015808 10226	implantation-associated protein	DKFZp564K1 42		NP_115497			11047007
C NM_178191 NP_835498 93974 ATPIFI ATPase inhil C NM_018434 NP_060904 55819 RNF130 ring finger pn C NM_004951 NP_060904 55819 RNF130 ring finger pn C NM_004951 NP_065090 1876 E2F6 E2F transcrip C NM_0020357 NP_065090 57092 PCNP PEST-contain C NM_006304 NP_997386 401388; FLJ42280 FLJ42280 pn C NM_006304 NP_997386 5797; 40256 FLJ42280 pn C NM_006923 NP_008854 6388 SDF2 stromal cell-d C NM_006923 NP_078805 79577 HRPT2 hyperparathyr NM_0024529 NP_001605 71 ACTG1 actin, gamma NM_001614 NP_001605 71 ACTG1 angiopoietin-l NM_005817 NP_003808 10226 M6PRBP1 mannose-6-ph NM_014236 NP_001422 2037 EPB41L2	protein phosphatase 1, catalytic subunit, beta iso	PPPICB		NP_996760			nioa8998
C NM_178191 NP_835498 93974 ATPIFI ATPase inhile C NM_018434 NP_060904 55819 RNF130 ring finger pn C NM_004951 NP_060904 55819 RNF130 ring finger pn C NM_004951 NP_065090 57092 PCNP E2F6 E2F transcrip C NM_020357 NP_065090 57092 PCNP PEST-contain C NM_006304 NP_997386 401388: 7979; FLJ42280 pr PCNP PEST-contain C NM_006304 NP_997386 401388: 7979; FLJ42280 pr PCNP PEST-contain C NM_006923 NP_620309 604;389185 BCL6 B-cell CLL/ly C NM_006923 NP_008854 6388 SDF2 stromal cell-d D NM_0024529 NP_001605 71 ACTG1 actin, gamma NM_001421 NP_001605 71 ACTG1 actin, gamma NM_001421 NP_001422 2037 EPB41L2	ribosomal protein S8	RPS8	6202; 94161	NP_001003	NM_001012	4	11046984
C NM_178191 NP_835498 93974 ATPIFI ATPase inhile C NM_018434 NP_060904 55819 RNF130 ring finger proposed in propo	glyceronephosphate O-acyltransferase	GNPAT		NP_055051	NM_014236		110289/0
C NM_178191 NP_835498 93974 ATPIFI ATPase inhile C NM_018434 NP_060904 55819 RNF130 ring finger proposed in propo	erythrocyte membrane protein band 4.1-like 2	EPB41L2		NP_001422	NM_001431	40	11028952
C NM_178191 NP_835498 93974 ATPIFI ATPase inhit C NM_018434 NP_060904 55819 RNF130 ring finger pp C NM_004951 NP_060904 1880 EB12 Epstein-Barr C NM_198258 NP_997705 1876 E2F6 E2F transcrip C NM_020357 NP_065090 57092 PCNP PEST-contain C NM_006304 NP_997386 401388; FLJ42280 FLJ42280 pro NM_006923 NP_620309 604; 389185 BCL6 B-cell CLL/ly NM_006923 NP_008854 6388 SDF2 stromal cell-d NM_001614 NP_078805 79577 HRPT2 hyperparathyr NM_004673 NP_001605 71 ACTG1 actin, gamma NM_012218 NP_114401 23731 C9orf3 chromosome s NM_012218 NP_703194 3609; ILF3 interleukin enl	mannose-6-phosphate receptor binding protein [M6PRBP1		NP_005808	NM_005817	40	11028946
C NM_178191 NP_835498 93974 ATPIFI ATPase inhit C NM_018434 NP_060904 55819 RNF130 ring finger pr C NM_004951 NP_060904 1880 EBI2 Epstein-Barr C NM_198258 NP_997705 1876 E2F6 E2F transcrip NM_020357 NP_065090 57092 PCNP PEST-contain NM_006304 NP_997386 401388; 402566 FLJ42280 FLJ42280 pro NM_006923 NP_620309 604; 389185 BCL6 B-cell CLL/by NM_006923 NP_008854 6388 SDF2 stromal cell-d NM_001614 NP_001605 71 ACTG1 actin, gamma NM_004673 NP_004664 9068 ANGPTL1 angiopoietin-l NM_032012 NP_114401 23731 C9orf5 chromosome 5	interleukin enhancer binding factor 3, 90kDa	ILF3	3609; 147727	NP_703194	NM_012218	40	11028912
C NM_178191 NP_835498 93974 ATPIFI ATPase inhit C NM_018434 NP_060904 55819 RNF130 ring finger pr C NM_004951 NP_064942 1880 EBI2 Epstein-Barr C NM_004951 NP_065090 57092 PCNP PEST-contain C NM_020357 NP_065090 57092 PCNP PEST-contain D NM_006304 NP_997386 401388; 402566 FLJ42280 FLJ42280 pr NM_006923 NP_620309 604; 389185 BCL6 B-cell CLL/ly NM_006923 NP_008854 6388 SDF2 stromal cell-d NM_001614 NP_001605 79577 HRPT2 hyperparathyr NM_004673 NP_004664 9068 ANGPTL1 angiopoietin-l	chromosome 9 open reading frame 5	C9orf5	23731	NP_114401	NM_032012	4C	11028886
C NM_178191 NP_835498 93974 ATPIFI ATPase inhit C NM_018434 NP_060904 55819 RNF130 ring finger pr C NM_004951 NP_060904 1880 EB12 Epstein-Barr C NM_198258 NP_997705 1876 E2F6 E2F transcrip NM_00304 NP_997386 401388; FLJ42280 FLJ42280 pro NM_006304 NP_997386 401388; FLJ42280 FLJ42280 pro NM_138931 NP_620309 604; 389185 BCL6 B-cell CLL/ly NM_006923 NP_008854 6388 SDF2 stromal cell-d NM_000204 NP_000195 3426 IF hyperparathyr NM_001614 NP_001605 71 ACTG1 actin, gamma	angiopoietin-like l	ANGPTLI	9068	NP_004664	NM_004673	†	nioa8864
C NM_178191 NP_835498 93974 ATPIFI ATPase inhit C NM_018434 NP_060904 55819 RNF130 ring finger pr C NM_004951 NP_004942 1880 EB12 Epstein-Barr C NM_198258 NP_997705 1876 E2F6 E2F transcrip NM_0020357 NP_065090 57092 PCNP PEST-contain NM_006304 NP_997386 401388; FLJ42280 FLJ42280 pr NM_138931 NP_620309 604; 389185 BCL6 B-cell CLL/ly NM_006923 NP_008854 6388 SDF2 stromal cell-d NM_0024529 NP_078805 79577 HRPT2 hyperparathyr NM_000204 NP_000195 3426 IF I factor (comp	actin, gamma l	ACTGI	71	NP_001605	NM_001614	£.	nioa8852
NM_178191 NP_835498 93974 ATPIFI ATPase inhit NM_018434 NP_060904 55819 RNF130 ring finger pr NM_004951 NP_004942 1880 EBI2 Epstein-Barr NM_198258 NP_997705 1876 E2F6 E2F transcrip NM_020357 NP_065090 57092 PCNP PEST-contain NM_006304 NP_997386 401388; 402566 FLJ42280 pr NM_138931 NP_620309 604; 389185 BCL6 B-cell CLL/ly NM_006923 NP_008854 6388 SDF2 stromal cell-d NM_024529 NP_078805 79577 HRPT2 hyperparathyr	I factor (complement)	IF	3426	NP_000195	NM_000204	40	nioassol
NM_178191 NP_835498 93974 ATPIFI ATPase inhit NM_018434 NP_060904 55819 RNF130 ring finger pr NM_004951 NP_004942 1880 EBI2 Epstein-Barr NM_198258 NP_997705 1876 E2F6 E2F transcrip NM_020357 NP_065090 57092 PCNP PEST-contain NM_06304 NP_997386 401388; 7979; 402566 FLJ42280 pr NM_138931 NP_620309 604; 389185 BCL6 B-cell CLL/ly NM_06923 NP_008854 6388 SDF2 stromal cell-d	hyperparathyroidism 2 (with jaw tumor)	HRPT2	79577	NP_078805	NM_024529	4C	nioa8811
NM_178191 NP_835498 93974 ATPIFI ATPase inhit NM_018434 NP_060904 55819 RNF130 ring finger pr NM_004951 NP_004942 1880 EBI2 Epstein-Barr NM_198258 NP_997705 1876 E2F6 E2F transcrip NM_020357 NP_065090 57092 PCNP PEST-contain NM_006304 NP_997386 401388; 7979; 402566 FLJ42280 pr NM_138931 NP_620309 604; 389185 BCL6 B-cell CLL/ly	stromal cell-derived factor 2	SDF2	6388	NP_008854	NM_006923	4C	nioa8796
NM_178191 NP_835498 93974 ATPIFI ATPase inhit NM_018434 NP_060904 55819 RNF130 ring finger pr NM_004951 NP_004942 1880 EBI2 Epstein-Barr NM_198258 NP_997705 1876 E2F6 E2F transcrip NM_020357 NP_065090 57092 PCNP PEST-contain NM_006304 NP_997386 401388; 402566 FLJ42280 pro	B-cell CLL/lymphoma 6 (zinc finger protein 51)	BCL6	604; 389185	NP_620309	NM_138931	4C	nioa8774
NM_178191 NP_835498 93974 ATPIFI ATPase inhit NM_018434 NP_060904 55819 RNF130 ring finger pr NM_004951 NP_004942 1880 EBI2 Epstein-Barr NM_198258 NP_997705 1876 E2F6 E2F transcrip NM_020357 NP_065090 57092 PCNP PEST-contair NM_006304 NP_997386 401388; FLJ42280 FLJ42280	,		7979; 402566				
NM_178191 NP_835498 93974 ATPIFI ATPase inhit NM_018434 NP_060904 55819 RNF130 ring finger pr NM_004951 NP_004942 1880 EBI2 Epstein-Barr NM_198258 NP_997705 1876 E2F6 E2F transcrip NM_020357 NP_065090 57092 PCNP PEST-contain	FLJ42280 protein	FLJ42280	401388;	NP_997386	NM_006304	8	mroa&//_
NM_178191 NP_835498 93974 ATPIFI ATPase inhit NM_018434 NP_060904 55819 RNF130 ring finger pr NM_004951 NP_004942 1880 EBI2 Epstein-Barr NM_198258 NP_997705 1876 E2F6 E2F transcrip	PEST-containing nuclear protein	PCNP	57092	NP_065090	MM_02035/		100000
NM_178191 NP_835498 93974 ATPIFI ATPase inhil NM_018434 NP_060904 55819 RNF130 ring finger pr NM_004951 NP_004942 1880 EBI2 Epstein-Barr	E2F transcription factor 6	E2F6	1876	NP_997705	8C7961 MINI	740	mioa864
NM_178191 NP_835498 93974 ATPIFI NM_018434 NP_060904 55819 RNF130	Epstein-Barr virus induced gene 2 (lymphocyte-spec	EBI2	1880	NP_004942	NM_004951	240	mioa854
NM_178191 NP_835498 93974 ATPIFI	ring finger protein 130	RNF130	55819	NP_060904	NM_018434	44C	mioa831
	ATPase inhibitory factor 1	ATPIFI	93974	NP_835498	NM_178191	2 4C	m10a760

RABIIA, member RAS oncogene family	RABIIA	8766	NP_004654	NM_004663	+	11100000//
valosin-containing protein	VCP	7415	NP_009057			miob000777 4C
splicing factor, arginine/serine-rich 2, interacti	SFRS2IP s		NP_004710		340	miob0496 4C
ubiquilin 2	UBQLN2	29978	NP_038472		140	miob0361 4C
hypothetical protein MGC33190	MGC33190 I	222255	NP_689962	NM_152749	74C	miob0207 4C
phosphoprotein enriched in astrocytes 15	PEA15	8682; 844	NP_003759	NM_003768	2/4C	miob0202 4C
Mlx interactor	MONDOA	22877	NP_055753	NM_014938	94C	m10b01894C
t-complex-associated-testis-expressed 1-like 1	TCTELI	6993	NP_006510		84C	m10b01/8/4C
hypothetical protein LOC51244	LOC51244	51244	NP_057558		7/4C	m10b01674C
PC326 protein	PC326	55827	NP_060912	NM_018442	440	m10b013414C
protein kinase C, nu	PRKCN	23683	NP_005804	NM_005813	340	m10a9935 4C
BTAF1 RNA polymerase II, B-TFIID transcription fac	BTAF1	9044	NP_003963	NM_003972		m10a9891 4C
STAM binding protein	STAMBP	10617	NP_998787	NM_006463	40	111089831140
topoisomerase (DNA) II alpha 170kDa	TOP2A	7153	NP_001058	NM_001067	240	min003/32 4C
chromosome 6 open reading frame 209	C6orf209	55788	NP_060838	NM_018368	46	micco702 4C
CocoaCrisp	LOC83690	83690	NP_113649	NM_031461	3 4C	m10a9666 4C
	KIAA0266	KIAA0266	9724	NM_021645	3 4C	m10a9049 4C
PMS1 postmeiotic segregation increased 1 (S. cerev	PMS1	5378	NP_000525	NM_000534	30 4C	m10a9630 4C
chaperonin containing TCP1, subunit 3 (gamma)	ССТЗ	7203	NP_005989	NM_005998	SI 4C	m10a9581 4C
zinc metalloproteinase (STE24 homolog, yeast)	ZMPSTE24	10269	NP_005848	NM_005857	40	m10a9510 4C
thioredoxin-like 2	TXNL2	10539	NP_006532	NM_006541	55 40	11104500340
CD109 antigen (Gov platelet alloantigens)	CD109	135228	NP_598000	NM_133493	73 40	mico0505 40
mitogen-activated protein kinase 9	МАРК9	5601	NP_620709	NM_139069	94 4C	mioa9294 4C
				NM_006585	79 4C	mioa9179 4C
hyaluronan and proteoglycan link protein	HAPLNI	1404	NP_001875	NM_001884	54 4C	mioa9154 4C

GNAS complex locus	CZAZ	2//8	111 00001	FINAL PARTY		
		3110	NP 536351	NM 080425	5 4C	miob2855 4C
cathepsin S	CTSS	1520	NP_004070	NM_004079	6 4 C	m1002836 4C
interferon gamma receptor 2 (interferon gamma tran	IFNGR2	3460	NP_005525	NM_005534	3 4 C	miob2743 4C
SNF2 histone linker PHD RING helicase	SHPRH	257218	NP_775105	NM_173082	5 4 C	miob2705/4C
salvador homolog 1 (Drosophila)	SAVI	60485	NP_068590	NM_021818	0/4C	miob2700 4C
N-glycanase 1	NGLYI	55768	NP_060767	NM_018297	7/4C	m10b2687 4C
G8 serologically defined colon cancer antigen 8	SDCCAG8	10806	NP_006633	NM_006642	2/4C	m10b2492 4C
H3 histone, family 3B (H3.3B)	нзғзв	3021	NP_005315	NM_005324	54C	m10b23/5/4C
Pirin	PIR	8544	NP_003653	NM_003662	40	m100222/4C
	NTANI	123803	NP_775745	NM_173474	57/4C	m10b2067/4C
glia maturation factor, gamma	GMFG	9535	NP_004868	NM_004877	8940	111001/89/4C
secreted phosphoprotein 1 (osteopontin, bone sialo	SPPI	6696	NP_000573	NM_000582	64 4C	m10b1 /34 4C
microfibrillar associated protein 5	MFAP5	8076	NP_003471	NM_003480)6/4C	miob1506 4C
				NM_014684)3 4C	m10b1493 4C
thrombospondin I	THBSI	7057	NP_003237	NM_003246	37 4C	miob1337 4C
DKFZP434F0 DKFZP434F091 protein	DKFZP 91	25917	NP_056268	NM_015453	20 40	111100132040
2 aarF domain containing kinase 2	ADCK2	90956	NP_443085	NM_052853	3140	miab1237 40
polycystic kidney disease 2 (autosomal dominant)	PKD2	5311	NP_000288	NM_000297	540	miob 1221 4C
acyl-CoA synthetase long-chain family member 4	ACSL4	2182	NP_075266	NM_022977	62/4C	m10b1062 4C
matrix Gla protein	MGP	4256	NP_000891	NM_000900	16 40	III10010104C
Al collagen, type III, alpha I (Ehlers-Danlos syndrom	COL3A1	1281	NP_000081	NM_000090	9140	misk 1017 4C
	RFXANK	8625	NP_604389	NM_003721	7440	
molecule possessing ankyrin repeats induced by lip	MAIL	64332	NP_113607	NM_031419	73 40	11110009/3/40
early endosome antigen 1, 162kD	EEAI	8411	NP_003557	NM_003566	39 4C	miob0939 4C
hypothetical protein FLJ10094	FLJ10094	55068	NP_060463	NM_017993	264C	miob0926 4C

1 1 2 2 2						
translocase of inner mitochondrial membrane 17 hom	TIMM17A transl	10440 TI	NP_006326	NM_006335	m10b4238 4C	m10b4
chromosome 20 open reading frame 104	C20orf104 chron	51230 C	NP_057520	NM_016436	m10b415/4C	niop4
CD151 antigen	CD151 CD1:	977 C	NP_620599	NM_004357	. 14152 4C	1004
paralemmin 2	PALM2 paral	114299; P. 11217	NP_443749	NM_00/203	miot4050 40	
HSPC154 protein	HSPC154 HSPO	29090 H	NP_054896	NM_014177	miob405 / 4C	
RNA binding motif (RNP1, RRM) protein 3	RBM3 RNA	5935 R	NP_006734	NM_006743	miob3982/4C	niob
ring finger protein 11	RNF11 ring	26994 R	NP_055187	NM_014372	miob3953 4C	niob
				XM_087525	miob3938 4C	niob
nuclear factor I/B	NFIB nucle	4781 N	NP_005587	NM_005596	m10b3911 4C	miob
S-phase kinase-associated protein 1A (p19A)	SKPIA S-ph	6500 S	NP_733779	NM_006930	m10b3898/4C	miob.
mitogen-activated protein kinase kinase kinase kin	MAP4K4 mito	9448 N	NP_663720	NM_145686	111003/63/4C	1100
Rho-associated, coiled-coil containing protein kin	ROCK1 Rho	6093 R	NP_005397	NM_005406	miob3696 4C	11100
metastasis suppressor l	MTSS1 meta	9788	NP_055566	NM_014751	miob3690/4C	11100
ubiquitin specific protease 47	USP47 ubiq	55031 L	NP_060414	NM_017944	1110b3684 4C	1100
signal transducing adaptor molecule (SH3 domain an	STAM sign	8027	NP_003464	NM_003473	misb3/8/1/4C	
non-metastatic cells 2, protein (NM23B) expressed	NME2 non-	4831	NP_002503	NM_002512	mich353114C	
low density lipoprotein receptor-related protein 8	LRP8 low	7804	NP_150643	NM_004631	miob34714C	
arginine-rich, mutated in early stage tumors	ARMET argi	7873	NP_006001	NM_006010	mich3471 4C	
Williams-Beuren syndrome chromosome region 1	WBSCR1 Wil	7458	NP_114381	NM_022170	miob3443 4C	1 00
WNT1 inducible signaling pathway protein 3	WISP3 WN	8838	NP_937882	NM_003880	miob3426/4C	100
ecotropic viral integration site 2A	EVI2A eco	2123	NP_055025	NM_014210	miob3411 4C) Jorun
				NM_002450	m10b3396 4C	mıot
heterogeneous nuclear ribonucleoprotein F	HNRPF hete	3185	NP_004957	NM_004966	miob3348 4C	miol
bone morphogenetic protein 6	BMP6 bon	654	NP_001709	NM_001718	miob3131 4C	miot
zinc finger protein 43 (HTF6)	ZNF43 zino	7594; 7757	NP_003414	NM_007153	m10b3044 4C	lorm

hypothetical protein BC018453	LOC129531 1	129531	NP_620153	NM_138798	· ·	1111000094 40
protein phosphatase 2, regulatory subunit B (B56).	PPP2R5C I	5527	NP_848703	NM_002719		miobe601/4C
activating signal cointegrator 1 complex subunit 1	ASCC1 2	51008	NP_057031	NM_015947) (.	miob860940
membrane-spanning 4-domains, subfamily A. member 6	MS4A6A	64231	NP_690591	NM_152852		m10b85/24C
F-box and leucine-rich repeat protein	FBXL11	22992	NP_036440	NM_012308		m1008341 4C
barrier to autointegration factor I	BANFI	8815	NP_003851	NM_003860		111008320140
zinc finger protein 331	ZNF331	55422	NP_061025	NM_018555		miob8214 4C
				NM_021243		IIII008143 4C
GTP binding protein overexpressed in skeletal musc	GEM	2669	NP_859053	NM_005261		111100/038/40
zinc finger, FYVE domain containing 21	ZFYVE21	79038	NP_076976	NM_024071		miob/391 4C
myotrophin	MTPN	136319	NP_665807	MM_145808		1,720
pleiotropic regulator 1 (PRL1homolog, Arabidopsis)	PLRG1	5356	NP_002660	NM_002669		mick7272 4C
DEAD (Asp-Glu-Ala-Asp) box polypeptide 3, X-linked	DDX3X	1654	NP_076829	NM_001356		mich7200 4C
eukaryotic translation initiation factor 2B. suhun	EIF2B1	1967	NP_001405	NM_001414		miob/109/40
transforming, acidic coiled-coil containing protei	TACC2	10579	NP_996744	NM_006997		mio 57100 40
cyclin B1 interacting protein 1	CCNB11P1	57820	NP_878272	NM_182851		111000090440
GABARAPL2 GABA(A) receptor-associated protein-like 2	GABARAPL2	11345	NP_009216	NM_00/285		miob6004 4C
protein associated with PRK1	AWP1	54469	NP_061879	NM 002002		miobósz
leucine rich repeat (in FLII) interacting protein	LRRFIPI	9208; 375316	NP_004726	NM_004735		mioh682114C
crystallin, alpha B	CRYAB	1410	NP_001876	NM_001885		minh6707 40
TBP-interacting protein	TIP120A	55832	NP_060918	NM_018448	AC 2	mioh6610
chemokine (C-X-C motif) ligand 9	CXCL9	4283	NP_002407	NM_002416	4C 4	miob6595 4C
eukaryotic translation initiation factor A	EIFIA	1964	NP_001403	NM_001412		mioh6562 40
DC2 protein	DC2	58505	NP_067050	NM_021227	4°C	miob6456 4C
KH domain containing, RNA binding, signal transduc	KHDRBSI	10657	NP_006550	NM_006559	40	miob6430/4C

					-	20102010VI
BCL2/adenovirus E1B 19kDa interacting protein 3-li	BNIP3L	665	NP_004322	NM_004331		111001170
XPA binding protein 1	XABI	11321	NP_009197	NM_00/266) (1001100 4
hypothetical protein FLJ11506	FLJ11506	79719	NP_078942	NM_024666		mioc1122 4C
hemicentin	FIBL-6	83872	NP_114141	NM_031935		mios1060 4C
Rho guanine nucleotide exchange factor (GEF) 17	ARHGEF17	9828	NP_055601	NM_014786	C	mioc0899 4C
				NM_020221	С	m10c0824 4C
upregulated during skeletal muscle growth 5	USMG5	84833	NP_116136	NM_032747	C	m10c0741 4C
transmembrane, prostate androgen induced RNA	TMEPAI	56937	NP_954640	NM_020182	Ċ	111100037340
phosphoglucomutase 2-like 1	PGM2L1	283209	NP_775853	NM_173582		min-0275 46
FLJ20073 protein	FLJ20073	54809	NP_060124	NM_017654		miaa0217 40
nuclear ubiquitous casein kinase and cyclin-depend	NUCKS	64710	NP_073568	NM_022/31		12103011
FK506 binding protein 5	FKBP5	2289; 401257	NF_004108	1111 OOT 11/	5	mioc0121 40
retinoblastoma binding protein 9	RBBP9	10741	NP_69/043	NM 004117		mioc0090 4C
proline arginine-rich end leucine-rich repeat prot	PRELP	5549	NP_958505	909900 WIN	10 6	miob9907/4C
neural precursor cell expressed, developmentally d	7 NEDD5	4735; 55917	NP_004395	NM_004404	10 1 to	miob9848 4C
DKFZp586G0 calcium-binding transporter	DKFZp580 123	29957	NP_998816	NM_013386		mish020240
fatty acid binding protein 1, liver	FABP1	2168	NP_001434	NM_001443	\$ 6	miob9284 4C
	KIAA1033	23325	NP_056090	NM_015275	5 6	111009183 40
	C14orf108	55745	NP_060699	NM_018229	4C	m10b9121 4C
Bardet-Biedl syndrome 2	BBS2	583	NP_114091	NM_031885	+6	11009003 40
membrane-spanning 4-domains, subfamily A, member 7	MS4A7	58475	NP_996823	NM_021201	† t	miob90524C
nascent-polypeptide-associated complex alpha polyn	' NACA	4666; 5557	NP_005585	NM_005594	40	miob8947/4C
cytochrome P450, family 1, subfamily B polymentia	CYPIBI	1545	NP_000095	NM_000104	40	m10b8932 4C
phosphonoformate immuno-associated protein 5	PFAAP5	10443	NP_055702	NM_014887	4C	m10b8812 4C
RAB14, member RAS oncogene family	RAB14	51552	NP_057406	NM_016322	4C	m10b8711 4C

N-myristoyltransferase 1	NMTI	4836	NF_000303			
RAB6A, member RAS oncogene family	RAB6A		NID OCCECE	NM 021079	††	mioc3716 4C
KAD4A, member RAS oncogene family		5070	NP 947500	NM 002869	4C	mioc3618 4C
PADAA	RAR4A	5867: 10638 RAR4A	NP_004569	NM_004578	4C	mioc3574 4C
olute carrier family 25 (miss t), octa potypepti	SLC25A3	5250	NP_998777	NM_002635	4C	m10c3430 4C
alcohol dehydrogenase [B (class I) beta nolymout:	ADHIB	125	NP_000659	NM_000668	40	IIII0C3208 4C
sodium channel modifier [SCNMI	79005	NP_076946	NM_024041	† †	min2200 4C
caldesmon I	CALDI	800	NP_149347	NM_033138	Ž d	mioc2706
nuclear domain 10 protein	NDP52	10241	NP_005822	NM_005831	40	mioc3127 4C
RAB5A, member RAS oncogene family	RAB5A	5868	NP_004153	NM_004162	5 5	mioc2950 4C
hypothetical protein MGC12466	MGC12466	93474	NP_149990	NM_033213	40	mioc2880/4C
ubiquitin specific protease 9 X-linked (for foot	USP9X	8239	NP_068706	NM_004652	40	200 40
PERP, TP53 apoptosis effector	PERP	64065	NP_071404	NM_022121	6	1111002/2040
peptidyl-prolyl isomerase G (cyclonhilin G)	PPIG	9360	NP_004783	NM_004792	1	
heterogeneous nuclear ribonucleoprofein H2 (H)	HNRPH2	3188	NP_062543	NM_019597	1	mina264 4C
TCDD-inducible poly(ADP-ribose) polymerase	TIPARP	25976	NP_056323	NM_015508	1	min 2502 40
zinc finger protein 265	ZNF265	9406	NP_976225	NM_005455	1	mio 252
proteasome (prosome, macropain) 26S subunit ATPac	PSMC2	5701	NP_002794	NM_002803		mioc2210 4C
protein kinase C-like 2	PRKCL2	5586	NP_006247	NIM 002502	340	mioc2173 4C
ARP8 actin-related protein 8 homolog (yeast)	ACTR8	93973	NP_075050	95C900 WIN	64C	mioc2166 4C
COX11 homolog, cytochrome c oxidase assembly prote	COXII	1353	NP_004366	NIM 000500	44C	mioc2074 4C
glioma tumor suppressor candidate region gene 2	GLTSCR2	29997	NP_056525	NM 004375	94C	mioc2039 4C
DKFZP564D116 protein	DKFZP564D1 16	26123	NP_056446	NIM 015510	040	mioc2019 4C
histone deacetylase 9	HDAC9	9734	NP_848512	NM 015631	8 4C	mioc 1928 4C
LIM domain containing preferred translocation part	LPP	4026	NP_005569	8/ CC00 MIN	9 4C	mioc1609 4C
				NIM ODEETO	10 40	mioc 1440 AC
	-			NM 005311)5 4C	mioc1205 4C

steroid sensitive gene 1	URB	151887	NP_955806	110661 199011		
cofactor required for Sp1 transcriptional activati	CRSP3	9439	NP_057063	NM 100511		mioc6260 4C
hypothetical protein MGC3222	MGC3222	79188	NP_077310	NM 004830		mioc6114 4C
actin, beta	ACTB	60; 71	NP_001092	NM_001101		mioc6075 4C
mucosa associated lymphoid tissue lymphoma translo	MALTI	10892	NP_776216	NM_006785		mioc 5740 4C
likely ortholog of mouse IRA1 protein	IRAI	79718	NP_078941	NM_024665		mioc3643 4C
tripartite motif-containing 16	TRIM16	10626	NP_006461	NM_006470		mio 27/12 4C
similar to RIKEN cDNA 1110018M03	LOC387758	387758	NP_976249	NM_2033/1		20000
twisted gastrulation homolog 1 (Drosophila)	TWSG1	57045	NP_065699	NM_020648	7	mioc5307 40
DKFZp564J157 protein	DKFZP564J1 57	54458	NP_060927	1010 OOC 10		mioc4925 4C
TATA box binding protein	TBP	6908	NP_003185	NM 018457	C	mioc4766 4C
uncharacterized hypothalamus protein HT011	Н1011	404636	NID 0000712	NM 003104	Ť	mioc4696 4C
palmitoyl-protein thioesterase 1 (ceroid-lipofusci	PILI	22025	NP 060942	NM 018472	†C	mioc4667 4C
ATP synthase, H+ transporting, mitochondrial F1 co	ATP5B	300	ND 0010//	NM 000310	†C	mioc4641 4C
actin related protein 2/3 complex, subunit 5, 16kD	ARPC5	10092	NF_005/08	NM 001686	4C	mioc4534 4C
polymerase (DNA directed), gamma	POLG	3428	ND 005700	NM 005717	4C	mioc4420 4C
emopamil binding protein-like	EBPL	64000	789C00 div	NM 002693	4C	mioc4366 4C
SM13 suppressor of mif two 3 homolog 2 (yeast)	2112117	01650	NP 115054	NM 032565	4C	mioc4318 4C
CDC 10 cell division cycle 10 homolog (S. cerevisia	CHCLMS	6613	NP 008868	NM_006937	4C	mioc4290 4C
CDC10 211 1: · · ·	CDC10	989	NP_001779	NM_001788	4C	mioc4247 4C
cell division cycle 26	CDC26	246184	NP_644815	NM_139286	4C	m10c4089 4C
hypothetical protein FLJ10970	FLJ10970	55273	NP_060756	NM_018286	#	770400111
core-binding factor, runt domain, alpha subunit 2.	CBFA2T1	862	NP_783554	INM_004349	Å t	niocana
serum response factor (c-fos serum response elemen	SRF	6722	NP_003122	NM_003131	À t	mioc3958 4C
muskelin I, intracellular mediator containing kelo	MKLNI	4289	NP_037387	NM_013255	40	mioc3906 4C
interleukin 18 binding protein	HL 18BP	10000	, , ,			2007

chromosome 6 open reading frame 111		25957	NP_116259	NM_032870	C	m10c84/9/4C
	FLJ37035 protein	FLJ3/035	377021	The state of the s		
dlhydropyrimidinase-like 2	DPYSL2	1000	200021	NM 207425	Č	mioc8434 4C
Convenie protein	David	1000	NP 001377	NM 001386	ţĊ	mioc8423 4C
MAWD hinding protein	MAWBP	64081	NP_071412	NM_022129	Ċ	mioc8410 4C
hypothetical protein BC008217	LOC92906	92906	NP_612403	NM_138394	†C	m10c85/9 4C
sel-1 suppressor of lin-12-like (C. elegans)	SELIL	6400	NP_005056	NM_005065		1111068007/40
transcriptional coactivator tubedown-100	TBDN100	80155	NP_476516	NM_057175		mioc/998 4C
Niemann-Pick disease, type C2	NPC2	10577	NP_006423	NM_006432	5 6	7000 40
hypothetical protein DC50	DC50	81892	NP_112487	NIN OILL	5 6	7005
hypothetical protein FLJ10986	FLJ10986	55277	NF_000/61	MM 031310	40	mioc7818 4C
chromosome 16 open reading frame 33	C16orf33	/9622	NP_0/884/	19C810 MN	40	mioc7807 4C
membrane protein, palmitoylated 6 (MAGUK p55 subfa	MPP6	51678	NP_057531	NM 024571	4C 4C	mioc7764 4C
uncharacterized hypothalamus protein HT007	HT007	55863	NP_060950	NM_018480	40	mioc7744 4C
prostaglandin E synthase	PTGES	9536	NP_942092	NM_004878	6	mino#7696 40
hypothetical protein FLJ22789	FLJ22789	84190	NP_115606	NM_032230	à t	mina75/1 1G
dishevelled associated activator of morphogenesis	DAAMI	23002	NP_055807	NM_014992	Å) &	mioc7500
protein tyrosine phosphatase, receptor type, C	PTPRC	5788	NP_563580	NM_002838	10 to	mioc7444 4C
chromosome 20 open reading frame 38	C20orf38	55304	NP_060797	NM_018327	6	mioc7270 4C
hypothetical protein FLJ20701	FLJ20701	55022	NP_060403	NM_017933	40	mioc/331/4C
kelch domain containing 2	KLHDC2	23588	NP_055130	NM_014315	1	. 222. 40
lymphocyte antigen 75	LY75	4065	NP_002340	INM_002349	3 8	mioc7760
DnaJ (Hsp40) homolog, subfamily B, member []	DNAJBII	51726	NF_05/390	NIM 002240	AC 6	mioc7170 AC
COP9 constitutive photomorphogenic homolog subunit	COPS5	10987	NP_006828	NM 016206	140	mioc6997 4C
chromosome 14 open reading frame 139	C14orf139	79686	NP_078909	NM_024633	740	mioc6987 4C
zinc finger, DHHC domain containing 7	ZDHHC7	55625	NP_060210	NM_017740	240	mioc6947 4C
				The state of the s		

mitochondrial ribosomal protein 1 50	MRPL50	54534	NP_061924	NM_019051	635 4C	miod2635/4C
protein kinase, cAMP-dependent, regulatory, type I	PRKARIA	5573	NP_997637	NM_002734	m10d2525/4C	miodz
translocase of inner mitochondrial membrane 17 hom	TIMM17A	10440	NP_006326	NM_006335	miod2412/4C	miod2
mastermind-like 2 (Drosophila)	MAML2	84441	NP_115803	NM_032427	m10d2330/4C	miodZ
solute carrier family 2 (facilitated glucose trans	SLC2A10	81031	NP_110404	NM_030777	miod2323/4C	miodz
5'-nucleotidase, ecto (CD73)	NT5E	4907	NP_002517	NM_002526	IIII002128 4C	L log
chromosome 10 open reading frame 118	C10orf118	55088	NP_060487	NM_018017	miod2025/4C	miodz
				NM_030921	m10d1942/4C	miodi
transforming growth factor beta 1 induced transcri	TGFB114	8848	NP_904358	NM_183422	miod1925/4C	miodi
MANSC domain containing 1	MANSCI	54682	NP_060520	NM_018050	miod1811/4C	miodi
protease, serine, 23	SPUVE	11098	NP_009104	NM_007173	IIII0d1558/4C	
hypothetical protein PRO1580	PRO1580	55374	NP_060972	NM_018502	miod1542/4C	Dorm
ribosomal protein S27-like	RPS27L	51065	NP_057004	NM_015920	IIII0011448/4C	
RABIIA, member RAS oncogene family	RABIIA	8766	NP_004654	NM_004663	miod1389 4C	miod
				XM_084964	m10d1331 4C	miod
hypothetical protein FLJ20445	FLJ20445	54708	NP_060294	NM_017824	miod1236/4C	miod
zinc finger protein 544	ZNF544	27300; 401932	NP_055295	NM_014480		
E2F transcription factor 3	E2F3	1871	NP_001940	NM_001949	miod1108 4C	miod
				XM_071675	miod0686 4C	miod
methylmalonyl Coenzyme A mutase	TUM	4594	NP_000246	NM_000255	m10d0592 4C	miod
Bardet-Biedl syndrome 7	BBS7	55212	NP_789794	NM_176824	m10d0441 4C	miod
hypothetical protein MGC8721	MGC8721	51669	NP_057211	NM_016127	miod0340/4C	miod
REV1-like (yeast)	REVIL	51455	NP_057400	NM_016316	m10c8694 4C	mioc
ribosomal protein L41	RPL41	6171	NP_066927	NM_021104	mioc8682 4C	mioc
myeloid cell leukemia sequence 1 (BCL2-related)	MCL1	4170	NP_877495	NM_021960	1111068019 40	mıoc

miod2639 4C	NM 006010	NP 006001	7873	ADMIT	
miod2665 4C	NM 033318	ND 201575	01/00	AKMEI	arginine-rich, mutated in early stage tumors
miod283714C	NIM 017087	7 2010/3	91089	LOC91689	hypothetical gene supported by AL449243
miod265/40	NM_01/987	NP_060457	55680	RUFY2	RUN and FYVE domain containing 2
miod35014C	NM_020536	NP_808779	57325	CSRP2BP	CSRP2 binding protein
11110d3391 4C	NM_004117	NP_004108	2289; 401257	FKBP5	FK506 binding protein 5
miod3854 4C	XM_032054		1011107		
miod4140 4C	NM_014781	NP 055596	1089	RRICCI	
miod4142.4C	024C10 WIN	- 000000	7021	KB ICCI	RB1-inducible coiled-coil 1
miod43424C	NM_012469	NP_036601	24148	C20orf14	chromosome 20 open reading frame 14
miod43484C	NM_001316	NP_803185	1434	CSEIL	CSE1 chromosome segregation 1-like (yeast)
miod+040740	NM_018048	NP_060518	55110	FLJ10292	hypothetical protein FLJ10292
miod440/4C	NM_002491	NP_002482	4709	NDUFB3	NADH dehydrogenase (ubiquinone) beta subcomplex
miod4449 4C	NM_080821	NP_543011	116151	C20orf108	chromosome 20 open reading frame 108
IIII004539/4C	NM_198057	NP_932174	1831	DSIPI	delta sleep inducing peptide, immunoreactor
	NM_172230	NP_757385	84447	HRD1	HRD1 protein
miod5030/4C	NM_007198	NP_009129	111212	PROSC	proline synthetase co-transcribed bomolog (box):
m10d512314C	NM_031431	NP_113619	83548	COG3	component of oligomeric galoi complay 3
miod5310/4C	NM_024579	NP_078855	79630	FLJ23221	hypothetical protein FI 173771
miod5349 4C	NM_152350	NP_689563	125144	MGC40157	hypothetical protein MCC40157
miod5369 4C	NM_020143	NP_064528	56902	LOC56902	putatative 28 kDa protein
miod5505 4C	NM_015310	NP_996792	23362	EFA6R	ADD ribosylation for
miod5651 4C	NM_018129	NP 060599		PNPO	nucleotide factor
miod5698 4C	NM_017935	NP 060405			Poll control oxidase
miod5703 4C	NM_004152	NP 004143			consisting death of the anklyrin repeats [
miod5707 4C	NM_003142				Siogram syndromy out: Program out of the syndromy out of the syndr
miod6029 4C	NM_052880	NP 443112	113791	MGC17330	HGET Company mingen in (automnigen La)
W. 12000.					Overs

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tumor differentially expressed 1	TDEI	10955	NP_945179	118900_141	t	
KIAA1287 protein	KIAA1287	57508	NP_065/99	NIM 000011	AC 6	
hyaluronan and proteoglycan link protein 1	HAPLNI	1404	NP_001875	NM_001884	40	ncr138/ ncr1437
				NM_014247	4C	ncr1352
NADH dehydrogenase (ubiquinone) beta subcomplex	NDUFB8	4714	NP_004995	NM_005004	4C	ncr1351
dolichyl-phosphate mannosyltransferase polypeptide	DPM1	8813	NP_003850	NM_003859	40	IICT 1 2 3 3
mesoderm induction early response 1	MI-ER1	57708	NP_065999	NM_020948	+	0/401011
integrin beta 3 binding protein (beta3-endonexin)	ІТGВЗВР	23421	NP_055103	NM_014288	4C	ncr0912
				NM_152446	4C	ncr0898
ribosomal protein L36a-like	RPL36AL	6166	NP_000992	NM_001001	4C	ncr0847
myotubular myopathy 1	MTMI	4534	NP_000243	NM_000252	4C	ncr0808
transcription factor Dp-2 (E2F dimerization partne	TFDP2	7029	NP_006277	NM_006286	#C	ncr0/66
beta-2-microglobulin	B2M	567	NP_004039	NM_004048	7	mc10/33
paraneoplastic antigen MA I	PNMAI	9240	NP_006020	NM_006029	à t	1100077
calmodulin I (phosphorylase kinase, delta)	CALMI	801	NP_008819	NM_006888	t t	mc=0624
putative nucleic acid binding protein RY-1	RY1	11017	NP_006848	NM_006857	\$ 6	710017
H3 histone, family 3B (H3.3B)	Н3F3B	3021	NP_005315	NM_005324	6 6	ncros4/
hypoxia-inducible protein 2	HIG2	29923	NP_037464	NM_013332	40	ncr0531
transferrin receptor (p90, CD71)	TFRC	7037	NP_003225	NM_003234	40	ncr04/8
SEC31-like 1 (S. cerevisiae)	SEC31L1	22872	NP_057295	NM_014953	i t	0.470
dynein, cytoplasmic, light intermediate polypeptid	DNCLII	51143	NP_057225	NM_016141	à t	mcr0/138
X-ray repair complementing defective repair in Chi	XRCC5	7520	NP_066964	NM_021141	† †	ncr0225
triple functional domain (PTPRF interacting)	TRIO	7204	NP_009049	NM_007118	4C	ncr0240
chromosome 10 open reading frame 10	C10orf10	11067	NP_008952	NM_007021	4C	ncr0238
RAB2, member RAS oncogene family	RAB2	5862	NP_002856	C98700 IMM	7	11010223

Williams Beuren syndrome chromosome region 20C	WBSCR20C	260294; 155400;	NP_684281	NM_145645	40	110100 4
acyl-CoA synthetase long-chain family member 4	ACSL4	2182	NP_075266	NM_022977		1
ribosomal protein L11	RPL11	6135	NP_000966	NM_0009/5		
tachykinin, precursor 1 (substance K, substance P,	TAC1	6863	NP_054704	NM_003182		
carnitine O-octanoyltransferase	CROT	54677	NP_066974	NM_021151	40	
chromosome 6 open reading frame 111	C6orf111	25957	NP_116259	NM_032870	40	
ring finger protein 149	RNF149	284996	NP_775918	NM_173647	40	
SKI-interacting protein	SNWI	22938	NP_036377	NM_012245	4C	
mitogen-activated protein kinase-activated protein	MAPKAPK5	8550	NP_620777	NM_139078	4C	
-				NM_004986	4C	
lung type-I cell membrane-associated glycoprotein	TIA-2	10630	NP_938203	NM_006474	4C	<u> </u>
ubiquitin specific protease 33	USP33	23032	NP_963920	NM_015017	4C	
TRAF family member-associated NFKB activator	TANK	10010	NP_597841	NM_004180	4C	
ATPase inhibitory factor I	ATPIF1	93974	NP_835498	NM_178191	4C	
forkhead box P1	FOXPI	27086	NP_116071	NM_032682	4C	
integral type I protein	P24B	23423	NP_031390	NM_007364	4C	
transmembrane, prostate androgen induced RNA	TMEPAI	56937	NP_954640	NM_020182	4C	
mitofusin l	MFNI	55669	NP_284941	NM_033540	4C	ncr2182
KIAA1536 protein	KIAA1536	57658	NP_065949	NM_020898	40	ncr2013
myosin, light polypeptide kinase	MYLK	4638	NP_444260	NM_053025	+	2012
A kinase (PRKA) anchor protein 13	AKAP13	11214	NP_658913	NM_055055	ĵ ŧ	nor1010
intracisternal A particle-promoted polypeptide	IPP	3652	NP_005888	NM_00/220	70 +6	ncr 1 876
hydroxysteroid (17-beta) dehydrogenase 7	HSD17B7	51478	NP_057455	NM_016371	7	nor1803
jumonji domain containing 1B	JMJDIB	51780	NP_057688	NM_016604	4C	ncr1563
				NM_014672	4C	ncr1550

4173 MCM4 MCM4 minichromosome maintenance deficient 4 (S. ce 2771 GNA12 guanine nucleotide binding protein (G protein), al 58508 MLL3 myeloid/lymphoid or mixed-lineage leukemia 3 10128 LRPPRC leucine-rich PPR-motif containing 11103 HRB2 HIV-1 rev binding protein 2 1173 AP2M1 adaptor-related protein complex 2, mu 1 subunit 23191 CYFIP1 cytoplasmic FMR1 interacting protein 1 5813 PURA purine-rich element binding protein A 3107 HLA-C major histocompatibility complex, class I, C 25949 P29 GCIP-interacting protein p29 10778 ZNF271 zinc finger protein 271 84707 BEX2 brain expressed X-linked 2 10972 TMP21 transmembrane trafficking protein 9510 ADAMTS1 a disintegrin-like and metalloprotease (reprolysin	NP_008919	NM_006988		L
MCM4 GNAI2 8 MLL3 8 LRPPRC 8 HRB2 AP2M1 CYFIP1 CYFIP1 CYFIP1 PURA PURA PLA-C HLA-C P29 ZNF271 ZNF271 BEX2 BEX2 B			17	1101/0/2
MCM4 GNAI2 BNAI3 MLL3 LRPPRC HRB2 AP2M1 CYFIP1 PURA PURA HLA-C HLA-C P29 ZNF271 ZNF271 BEX2	NP_006818	178900 FAIN	5 8	
MCM4 GNAI2 RAPPRC HRB2 AP2MI CYFIPI CYFIPI HLA-C	NP_116010	NM_032621	40	ncr7668
MCM4 GNAI2 RMLL3 RHRB2 AP2MI CYFIPI CYFIPI PURA HLA-C HLA-C P29	NP_006620	NM_006629	70 40	
MCM4 GNAI2 RMLL3 REPPRC HRB2 AP2MI CYFIPI CYFIPI PURA HLA-C HLA-C	NP_997053	NM_015484	10 40	ncr7245
MCM4 GNAI2 8 MLL3 8 LRPPRC 3 HRB2 AP2MI 1 CYFIPI PURA	NP_002108	NM_002117	46	1007 TOOL
MCM4 GNAI2 8 MLL3 8 LRPPRC 3 HRB2 AP2MI 1 CYFIPI	NP_005850	NM_00215	<u> </u>	ncr7082
MCM4 GNAI2 MLL3 LRPPRC HRB2 AP2M1	NP_055423	NM_014608	7	ncr6335
MCM4 GNAI2 MLL3 LRPPRC HRB2	NP_004059	NM_004068	<u> </u>	ner6333
MCM4 GNAI2 MLL3 LRPPRC	NP_008974	NM_007043	t t	ncr6143
MCM4 GNAI2 MLL3	NP_573566	NM_133259	40 40	ncr61/1
MCM4 GNAI2	NP_733751	NM_1/0606	j t	nor6077
MCM4	NP_002061	NM_002070	À t	ncr5020
1 1 1 1 1 1 1 1 1 1 1 1 1	NP_877423	NM_005914	\$ 6	mar 5713
9100 USP10 ubiquitin specific protease 10	NP_005144	NM_005153	5 6	11013400
51366 DD5 progestin induced protein	NP_056986	NM_015902	à t	nou.\$ 100
3107 HLA-C major histocompatibility complex, class I, C	NP_002108	NM_002117	à t	ncr\$160
7716 ZNF161 zinc finger protein 161	NP_009077	NM_00/146	5 7	mcr5066
84271 POLDIP3 polymerase (DNA-directed), delta interacting prote	NP_835237	NM_032311	20 6	ncr4656
		XM_301731	5	11014330
163859 FLJ35382 hypothetical protein FLJ35382	NP_689821	8097C1 WN	à đ	204041011
23404 EXOSC2 exosome component 2	NP_055100	NM_014285	3 8	ncr/15/15
2199 FBLN2 fibulin 2	NP_001989	NM_001998	Ž t	ncr/1520
1499 CTNNB1 catenin (cadherin-associated protein), beta 1, 88k	NP_001895	NM_001904	1 C	ncr4416
201134 MGC33887 hypothetical protein MGC33887	NP_659473	NM_145036	45	IICI4202

complement component 1, q subcomponent binding pro	CIQBP	708	NP_001203	NM_001212	40	nerb0033
retinoblastoma binding protein 6	RBBP6	5930	NP_116015	NM_006910	4C	
RAB, member RAS oncogene family-like 5	RABL5	64792	NP_073614	NM_022777	4C	1
UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltr	B3GNT6	11041	NP_006867	NM_006876	4C	
solute carrier family 39 (zinc transporter), membe	SLC39A1	27173	NP_055252	NM_014437	4C	
zinc finger protein 36, C3H type, homolog (mouse)	ZFP36	7538	NP_003398	NM_003407	4C	
superoxide dismutase 2, mitochondrial	SOD2	6648	NP_000627	NM_000636	4C	
HSPC182 protein	HSPC182	29101	NP_054907	NM_014188	4C	
				NM_153366	4C	
ubiquitin-conjugating enzyme E2L 3	UBE2L3	7332	NP_937800	NM_003347	4C	
K+ channel tetramerization protein	GMRP-1	84280	NP_115696	NM_032320	4C	ncr8893
serine (or cysteine) proteinase inhibitor, clade B	SERPINB9	5272	NP_004146	NM_004155	1 6	
inhibitor of kappa light polypeptide gene enhancer in B-cells, kinase complex-associated protein	IKBKAP	8518; 10880 IKBKAP	NP_003631	NM_003640	40	
COMM domain containing 10	COMMD10	51397	NP_057228	NM_016144	40	
histone I, H4f	HIST1H4F	8361; 8364	NP_003531	NM_003542	4C	ncr8693
bromodomain containing 2	BRID2	6046	NP_005095	NM_005104	4C	ncr8588
basic transcription factor 3	BTF3	689; 220793 BTF3	NP_001198	NM_001207	4C	ncr8542
cholinergic receptor, muscarinic 3	CHRM3	1131	NP_000731	NM_000740	4C	ncr8290
ornithine decarboxylase antizyme inhibitor	OAZIN	51582	NP_680479	NM_015878	4C	ncr8067
ADP-ribosylation factor-like 5	ARL5	26225	NP_817114	NM_012097	4C	ncr8041
DC2 protein	DC2	58505	NP_067050	NM_021227	40	ncr/9/3
tyrosine 3-monooxygenase/tryptophan 5-monooxygenas	YWHAG	7532	NP_036611	NM_012479	40	ncr/904
mutL homolog 1, colon cancer, nonpolyposis type 2	MLHI	4292	NP_000240	NM_000249	4C	ncr7852
FSHD region gene 1	FRG1	2483	NP_004468	NM_004477	4C	ncr7813
hippocampus abundant transcript	HIATI	64645	NP_149044	NM_033055	4C	ncr7768

4 8106 4 8106 9 57690 7 80267 2 26057 2 26057 2 36910 3 56910 3 56910 3 56910 5 56910 5 56910	NP_001604 NP_112506 NP_060516	NM 014892	\$	ncrb3942
4 8106 PABPNI 9 57690 KIAA1582 7 80267 Clorf22 2 26057 ANKRD17 6 4836 NMT1 7 56910 STARD7 7071 TIEG 10616 C20orf18	NP_001604 NP_112506 NP_060516	0400 IVIVI		
4 8106 PABPNI 9 57690 KIAA1582 7 80267 C1orf22 2 26057 ANKRD17 8 4836 NMT1 5 56910 STARD7 7071 TIEG 10616 C20orf18	NP_001604 NP_112506	NIM DIODAL	4C	ncrb3936
1 8106 PABPNI 57690 KIAA1582 80267 C1orf22 26057 ANKRD17 4836 NMT1 56910 STARD7 7071 TIEG	NP_001604	NM_031229	4C	ncrb3702
8106 PABPNI 57690 KIAA1582 80267 C1orf22 26057 ANKRD17 4836 NMT1 56910 STARD7 7071 TIEG		NM_001613	4C	
8106 PABPNI 57690 KIAA1582 80267 C1orf22 26057 ANKRD17 4836 NMT1 56910 STARD7	NP_005646	NM_005655	4°C	
8106 PABPN1 57690 KIAA1582 80267 C1orf22 26057 ANKRD17	NP_644672	NM_020151	40	ncrb3301 4C
8106 PABPN1 57690 KIAA1582 80267 C1orf22 26057 ANKRD17	NP_066565	NM_021079	4C	ncrb3284 4C
8106 PABPNI 57690 KIAA1582 80267 C1orf22	NP_942592	NM_032217	40	nerb3003
8106 PABPNI 57690 KIAA1582	NP_079467	NM_025191	4C	
8106 PABPNI	NP_061869	NM_018996	4C	
	NP_004634	NM_004643	4°C	ncrb2288
81631 MAPILC3B	NP_073729	NM_022818	16	nero2266 4C
84 3122 HLA-DRA major histocompatibility complex, class II, DR alp	NP_061984	NM_019111	40	nerb2092 4C
54466 SPIN2	NP_061876	NM_019003	40	10101936 4C
51 8672 EIF4G3 eukaryotic translation initiation factor 4 gamma,	NP_003751	NM_003760	4C	ncrb1802 4C
		NM_031407	4C	ncrb1456 4C
90441 ZNF622	NP_219482	NM_033414	4C	ncrb1438 4C
384 ARG2	NP_001163	NM_001172	4C	ncrb1373
629 BF	NP_001701	NM_001710	4C	ncrb0782 4C
9733 SART3	NP_055521	NM_014706	16	ncrouo96
NAD(P)H dehydrogenase, quinone l	NP_000894	NM_000903	1	100013
3134 HLA-F	NP_061823	NM_018950		11C100328
128 ADH5 alcohol dehyd	NP_000662	NM_000671		ncrb0262
29082 C14orf123	NP_054888	NM_014169		ncrb0145
031 521 ATP5I ATP synthase, H+ transporting, mitochondrial F0 co	NP_009031	NM_007100	14°C	ncrb0054 4C

NP_073741 64852 RBM21 RNA binding motif protein 21			
	NM_022830	100	10001
•	NM 022830	8 4C	ncrb6818
NP_996813 54843 SYTL2 synaptotagmin-like 2	NM_032943	1 4C	ncrb6661 4C
NP_001322 1500 CTNND1 catenin (cadherin-associated protein), delta	NM_001331	0 4C	ncrb6530
NP_064518 56895; AGPAT4 1-acylglycero1-3-phosphate O-acyltransferase 4 (ly	NM_020133	3 4C	ncrb6453
NP_116021 10914 PAPOLA poly(A) polymerase alpha	NM_032632		ncrb6432
NP_001688 539 ATP5O ATP synthase, H+ transporting, mitochondrial F1 co	NM_001697		ncrb6327
	NM_032754	2 4C	ncrb6192 4C
NP_057046 51077 C14orf111 chromosome 14 open reading frame 111	NM_015962	7 4C	ncrb6087 4C
NP_001531 3315 HSPB1 heat shock 27kDa protein 1	NM_001540	7 4C	ncrb6037 4C
NP_002309 4017 LOXL2 lysyl oxidase-like 2	NM_002318	5 4C	ncrb5595
NP_066972 23406 COTL1 coactosin-like I (Dictyostelium)	NM_021149	77 4C	ncrb5197
NP_003872 8839 WISP2 WNT1 inducible signaling pathway protein 2	NM_003881	1	ncrb4957
NP_003609 8491 MAP4K3 mitogen-activated protein kinase kin	NM_003618	2 4C	ncrb4912 4C
NP_116787 839 CASP6 caspase 6, apoptosis-related cysteine protease	NM_001226	8 4C	ncrb4538 4C
NP_004538 4710 NDUFB4 NADH dehydrogenase (ubiquinone) beta subcomplex,	NM_004547	40	ncrb4441 4C
NP_000963 6130; RPL7A ribosomal protein L7a	NM_000972	90 4C	ncrb4390 4C
	XM_211709	39 4C	ncrb4339
NP_699196 202018 FLJ90013 hypothetical protein FLJ90013	NM_153365	19 4C	ncrb4319
NP_061060 55454 GALNACT-2 chondroitin sulfate GalNAcT-2	NM_018590	82 4C	ncrb4182
NP_851824 122704 MRPL52 mitochondrial ribosomal protein L52	NM_181304	39 4C	ncrb4039 4C

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eukaryotic translation elongation factor alpha	EEF1A1 eı	1915; 389411	NP_001393	NM_001402	40	ncrc0239 40
geranylgeranyl diphosphate synthase l	GGPS1 g	9453	NP_004828	NM_004837	4C	nerc01/4
beta-2-microglobulin	B2M b	567	NP_004039	NM_004048	40	
AT rich interactive domain 5B (MRF1-like)	ARID5B A	84159		XM_084482	4C	1
CocoaCrisp	LOC83690 C	83690	NP_113649	NM_031461	4C	ı
expressed in T-cells and eosinophils in atopic der	ETEA e	23197	NP_055428	NM_014613	4C	1
solute carrier family 25 (mitochondrial carrier; p	SLC25A3 s	5250	NP_998777	NM_002635	4C	
hypothetical protein FLJ10849	FLJ10849 h	55752	NP_060713	NM_018243	4C	ncrb8820 4C
ribonuclease P 21kDa subunit	RPP21 r	79897	NP_079115	NM_024839	4C	ncrb8790 4C
formin-like 2	FMNL2 f	114793	NP_443137	NM_052905	40	ncrb8/51/4C
kinesin family member 3B	KIF3B k	9371	NP_004789	NM_004798	40	ncrb8693
delta sleep inducing peptide, immunoreactor	DSIPI c	1831	NP_932174	NM_198057	40	ncrb8665
CGI-125 protein	CGI-125	51003	NP_057144	NM_016060		ncrb8649
	synaptopodin 2	SYNPO2	171024	XM_050219		IICI08451
eukaryotic translation elongation factor alpha	EEF1A1 6	1915; 389411	NP_001393	NM_001402	40	ncro843 / 40
S100 calcium binding protein A6 (calcyclin)	S100A6	6277; 400825	NP_055439	NM_014624	6	ncrb8392
collagen, type X, alpha 1(Schmid metaphyseal chond	COL10A1	1300	NP_000484	NM_000493	4C	ncrb8385
tensin-like SH2 domain containing 1	TENSI	64759	NP_073585	NM_022748		ncrb8343
ADP-ribosylation factor GTPase activating protein	ARFGAP3	26286	NP_055385	NM_014570		ncrb8273
nitrilase I	ITIN	4817	NP_005591	NM_005600		ncrb8253
acetoacetyl-CoA synthetase	AACS	65985	NP_076417	NM_023928		ncrb8239
phosphatidic acid phosphatase type 2B	PPAP2B	8613	NP_803133	NM_003713	4C	ncrb8207 4C
				NM_021826	4C	ncrb8203 4C
chromosome 10 open reading frame 104	C10orf104	119504	NP_775744	NM_173473	1 4C	ncrb8134 4C

A I P synthase, H+ transporting, mitochondrial F0 co	AIPJ	1	141 0010/0		,	
RAD21 homolog (S. pombe)	RAD21		NP_006236	NM 000500	40	nere1751 /
ankylosis, progressive homolog (mouse)	ANKH	2	NP_473368	NM_054027	40 6	nere 1 505
solute carrier family 30 (zinc transporter), membe	SLC30A5		NP_076960	NM_022902	5 6	
lysosomal-associated protein transmembrane 4 alpha	LAPTM4A	9741	NP_055528	NM_014713	4C	1
RAB10, member RAS oncogene family	RAB10	10890	NP_057215	NM_016131	4C	1
ATP-dependent glucokinase	ADP-GK	83440	NP_112574	NM_031284	4C	
ras homolog gene family, member B	RHOB	388	NP_004031	NM_004040	4C	
myristoylated alanine-rich protein kinase C substr	MARCKS	4082	NP_002347	NM_002356	4C	1
thioredoxin interacting protein	TXNIP	10628	NP_006463	NM_006472	4C	
ubiquitination factor E4B (UFD2 homolog, yeast)	UBE4B	10277	NP_006039	NM_006048	4C	1
ring-box 1	RBX1	9978	NP_055063	NM_014248	4C	ncrc0846 4C
proteasome (prosome, macropain) subunit, alpha typ	PSMAI	5682	NP_683877	NM_148976	4C	ncrc0803 4C
integrin, beta-like 1 (with EGF-like repeat domain	ITGBL1	9358	NP_004782	NM_004791	4C	ncrc0/93
RALBPI associated Eps domain containing 1	REPS1	85021	NP_114128	NM_031922	#C	ncrc0/29/40
mitogen-activated protein kinase kinase kinase 4	MAP3K4	4216	NP_006715	NM_005922	4C	ncrc0715 4C
collagen, type V, alpha 2	COL5A2	1290	NP_000384	NM_000393	4C	ncrc0/04 4C
hypothetical protein, clone 2746033	HSA272196	55352	NP_060875	NM_018405	4C	ncrc0667
2-hydroxyphytanoyl-CoA lyase	HPCL2	26061	NP_036392	NM_012260	4C	
suppression of tumorigenicity 13 (colon carcinoma)	ST13	6767	NP_003923	NM_003932	4C	
DnaJ (Hsp40) homolog, subfamily B, member 9	DNAJB9	4189	NP_036460	NM_012328	40	nere0413
proteasome (prosome, macropain) subunit, beta type	PSMB4	5692	NP_002787	NM_002796	100	nereosas
scavenger receptor class F, member 1	SCARF1	8578; 83547	NP_663327	NM_145351	† C	ncrc0342
BTG3 associated nuclear protein	BANP	54971	NP_524576	NM_017869	4C	ncrc0304 4C
phenylalanine-tRNA synthetase-like, beta subunit	FARSLB	10056	NP_005678	/ 000000 /	i	110100272

interleukin 1 receptor accessory protein	ILIRAP	3556	NP_608273	NM_002182	40	ncrc3434
DEAD (Asp-Glu-Ala-Asp) box polypeptide 24	DDX24	57062	NP_065147	NM_020414	4C	ncrc3416
S-phase kinase-associated protein 1A (p19A)	SKP1A 9	6500	NP_733779	NM_006930	4C	ncrc3358 4
apolipoprotein A-I binding protein	APOA1BP :	128240	NP_658985	NM_144772	4C	ncrc3276 '
bobby sox homolog (Drosophila)	ВВХ	56987	NP_064620	NM_020235	4C	ncrc3258
integral membrane protein 2B	ITM2B	9445	NP_068839	NM_021999	4C	
SEC31-like I (S. cerevisiae)	SEC31L1	22872	NP_057295	NM_014933	†C	nere3072 4C
FK506 binding protein 14, 22 kDa	FKBP14	55033	NP_060416	NM_017946	4C	ncrc3045
Bardet-Biedl syndrome 7	BBS7	55212	NP_789794	NM_176824	4C	ncrc2940 4C
sarcolipin	SLN	6588	NP_003054	NM_003063	4C	ncrc2928
hypothetical protein MGC2747	MGC2747	79086	NP_077009	NM_024104	4C	.i.
pleckstrin homology domain interacting protein	PHIP	55023	NP_060404	NM_017934	4C	1
influenza virus NS1A binding protein	IVNSIABP	10625	NP_006460	NM_006469	4C	1
proteasome (prosome, macropain) subunit, alpha typ	PSMA5	5686	NP_002781	NM_002790	4C	1
retinoic acid induced 16	RAII6	64760; 55307	NP_073586	NM_022749	4C	
KIAA0433 protein	KIAA0433	23262	NP_056031	NM_015216	4C	1
hypothetical protein FLJ10637	FLJ10637	55726	NP_060634	NM_018164	4C	1
eukaryotic translation initiation factor 3 subunit	elF3k	27335	NP_037366	NM_013234	4C	ncrc2273
decorin	DCN	1634	NP_598014	NM_001920	4C	ncrc2119
Rho GTPase activating protein 6	ARHGAP6	395	NP_038286	NM_001174	4C	ncrc2018
interferon induced transmembrane protein 3 (1-8U)	IFITM3	10410	NP_066362	NM_021034	4C	ncrc1999
multiple coagulation factor deficiency 2	MCFD2	90411	NP_644808	NM_139279	4C	
dUTP pyrophosphatase	DUT	1854	NP_001939	NM_001948	4C	
actin, alpha 2, smooth muscle, aorta	ACTA2	59	NP_001604	NM_001613	4C	ncrc1873 4C
TCDD-inducible poly(ADP-ribose) polymerase	TIPARP	25976	NP_056323	NM_015508	4C	ncrc1765

transient receptor potential cation channel, subfa	TRPC4AP	26133	NP_955400	NM_015638	4C	ncrc4808
follistatin-like 1	FSTL1	11167	NP_009016	NM_007085	4C	ncrc4794
chromosome 14 open reading frame 4	C14orf4	64207	NP_078772	NM_024496	4C	ncrc4757
zinc finger, ZZ-type with EF hand domain 1	ZZEF1	23140	NP_055928	NM_015113	4C	ncrc4654
protein tyrosine phosphatase, non-receptor type 12	PTPN12	5782	NP_002826	NM_002835	4C	ncrc4633
				NM_014739	tC Ot	ncrc4600 4C
homocysteine-inducible, endoplasmic reticulum stre	HERPUDI	9709	NP_055500	NM_014685	4C	ncrc4597
carboxypeptidase A3 (mast cell)	CPA3	1359	NP_001861	NM_001870	4C	ncrc4531
vacuolar protein sorting 54 (yeast)	VPS54	51542	NP_057600	NM_016516	4C	ncrc4313
helicase, ATP binding I	HELICI	10973; 63921	NP_006819	NM_006828	4C	ncrc4296
amyloid beta (A4) precursor protein-binding, famil	APBBIIP	54518	NP_061916	NM_019043	4C	ncrc4247
platelet-derived growth factor receptor-like	PDGFRL	5157	NP_006198	NM_006207	4C	ncrc4226
translocase of inner mitochondrial membrane 17 hom	TIMM17A	10440	NP_006326	NM_006335	4C	ncrc4089
phosphatidic acid phosphatase type 2B	PPAP2B	8613	NP_803133	NM_003713	4C	ncrc4079
ubiquitin-like 3	UBL3	5412	NP_009037	NM_007106	4C	ncrc4047
nuclear ubiquitous casein kinase and cyclin-depend	NUCKS	64710	NP_073568	NM_022731	4C	ncrc4000
eukaryotic translation initiation factor 3, subuni	EIF3S5	8665	NP_003745	NM_003754	4C	ncrc3953
small fragment nuclease	DKFZP566E1 44	25996	NP_056338	NM_015523	4C	ncrc3840
HSPC056 protein	HSPC056	25852	NP_998819	NM_014154	4C	ncrc3624
ribosomal protein L6	RPL6	6128	NP_000961	NM_000970	4C	ncrc3544
FLJ20298 protein	FLJ20298	54885	NP_942582	NM_017752	4C	ncrc3536
phosphoglycerate mutase 1 (brain)	PGAMI	5223	NP_002620	NM_002629	4C	ncrc3529
ATP synthase, H+ transporting, mitochondrial F0 co	ATP5J	522	NP_001676	NM_001685	4C	ncrc3468
heme binding protein 2	HEBP2	23593	NP_055135	NM_014320	4C	ncrc3464
glutaminase	GLS	2744	NP_055720	NM_014905	4C	ncrc3453
		4	·			

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icity 13 (colon carcinoma)	suppression of tumorigenicity 13 (colon carcinoma)	ST13	6767	NP_003923	ININI_003932		
nt binding protein 1	cAMP responsive element binding protein	CREB1	1385	NP_604391	NM_134442		
ncogene family	RAB9A, member RAS oncogene family	RAB9A	9367	NP_004242	NM_004251		nere5569 40
	AXIN1 up-regulated 1	AXUD1	64651	NP_149016	NM_033027	<u>†</u>	
ı (syntenin)	syndecan binding protein (syntenin)	SDCBP	6386	NP_005616	NM_005625	† †	. 1
group nucleosome binding domain 1	high-mobility group nuc	HMGN1	3150	NP_004956	NM_004965	6	
	golgi apparatus protein	GLG1	2734	NP_036333	NM_012201	40	
Jumonji, AT rich interactive domain 1B (RBP2-like)	Jumonji, AT rich interac	JARIDIB	10765	NP_006609	NM_006618	4C	1
of Charlevoix-Saguenay (sacsin)	spastic ataxia of Charley	SACS	26278	NP_055178	NM_014363	t	
	zinc finger protein 577	ZNF577	84765	NP_116068	NM_032679		10103207 4C
amily member 6	transmembrane 4 superfamily member 6	TM4SF6	7105	NP_003261	NM_003270	4	20103101
membrane protein, palmitoylated 6 (MAGUK p55 subfa	membrane protein, paln	MPP6	51678	NP_057531	NM_016447	#6	
	matrilin 3	MATN3	4148	NP_002372	NM_002381		100001
itiation factor 3, subuni	eukaryotic translation initiation factor 3, subuni	EIF3S6	3646	NP_001559	NM_001568	10 to	ncrc5001 4C
eukaryotic translation initiation factor 3 subunit	eukaryotic translation ii	eIF3k	27335	NP_037366	NM_013234	5 8	I
translocase of outer mitochondrial membrane 70 hom	translocase of outer mit	ТОММ70А	9868	NP_055635	NM_014820	5 6	nereso72
	senataxin	KIAA0625	23064	NP_055861	NM_015046	\$ 6	
J13352	hypothetical protein FLJ13352	FLJ13352	79644	NP_078868	NM_024592	4C	
link domain containing 1	extracellular link doma	XLKDI	10894	NP_006682	NM_006691	4C	
	opioid receptor, sigma	OPRS1	10280	NP_671516	NM_147160	+	
	CDC-like kinase 1	CLK1	1195	NP_004062	NM_0040/1	40 8	
factor XIII, A1 polypeptide	coagulation factor XIII	F13A1	2162	NP_000120	NIM 004021	40	
protein	PAI-1 mRNA-binding protein	PAI-RBP1	26135	NP_056455	NM 000120	4C	ncrc4985
ATP synthase, H+ transporting, mitochondrial F1 co	ATP synthase, H+ trar	ATP5E	514	NP_008817	NM_006886	7 6	ncrc4885
4	formin binding protein 4	FNBP4	23360	NP_036123	010000		

8 4C NM_005013 NP_005004 4925; 399873 11 4C NM_017924 NP_060394 55017 33 4C NM_015411 NP_060394 55017 8 4C NM_05176 NP_005167 517; 34141; 8 4C NM_024078 NP_076983 79050 0 4C NM_021222 NP_067045 58497; 14 4C NM_003713 NP_803133 8613 5 4C NM_003713 NP_803133 38613 5 4C NM_002247 NP_002238 3778 6 4C NM_003518 NP_002957 6281 7 4C NM_000581 NP_0055744 22868 3 4C NM_007081 NP_005744 22868 3 4C NM_007081 NP_0057076 966 NP_007076 NP_007676 966 NP_007676 NP_0057659 51559 4C NM_003418 NP_003406 10959 4C NM_0004487 NP_004478 2804 4C NM_0005801 NP_00	chromosome 6 open reading frame 62	C6ort62 c	81088	NF_112201	1011_00000		1
8 4C NM_005013 NP_005004 4925; 39873 NUCB2 11 4C NM_017924 NP_060394 55017 C14orf119 33 4C NM_015411 NP_060394 55017 C14orf119 8 4C NM_015411 NP_056226 25870 SUMF2 2 4C NM_024078 NP_067045 58497; HTCD37 4 4C NM_025266 NP_067045 58497; HTCD37 4 4C NM_003713 NP_803133 8613 PPAP2B 5 4C NM_002247 NP_002238 3778 KCNMA1 7 4C NM_002966 NP_002957 6281 S100A10 1 4C NM_002966 NP_002957 6281 S100A10 2 4C NM_002909 NP_055744 22868 KIAA0971 3 4C NM_003345 NP_0037417 29889 HUMAUANT 4 C NM_003455 NP_037417 29889 HUMAUANT 4 C NM_003455 NP_03606 10959 RNP20 4 C <td< td=""><td>outative translation initiation factor</td><td></td><td></td><td>ND 112201</td><td>020020 MN</td><td>‡C</td><td></td></td<>	outative translation initiation factor			ND 112201	020020 MN	‡C	
4C NM_005013 NP_005004 4925: 39873 NUCB2 11 4C NM_017924 NP_060394 55017 C14orf119 33 4C NM_017924 NP_060394 55017 C14orf119 8 4C NM_005176 NP_076983 79050 SUMF2 0 4C NM_024078 NP_076983 79050 MGC3162 4 4C NM_003506 NP_067045 58497: HTCD37 5 4C NM_003713 NP_803133 8613 PPAP2B 5 4C NM_003713 NP_002238 3778 KCNMA1 7 4C NM_003518 NP_002957 6281 S100A10 3043 1 4C NM_003518 NP_0055744 22868 KIAA0971 NP_005676 2 4C NM_0014929 NP_055744 22868 KIAA0971 NP_0057676 3 4C NM_0014929 NP_055744 22868 KIAA0971 NP_005676 4 C NM_001482 NP_0057676 966 CD59 HUMAUAUT 5 4C NM_003455	of the state of th			NP 005792	NM 005801	†	ncrc8841
4C NM_005013 NP_005004 4925; 399873 NUCB2 11 4C NM_017924 NP_060394 55017 C14orf119 33 4C NM_017921 NP_060394 55017 C14orf119 8 4C NM_024078 NP_076983 79050 MGC3162 6 4C NM_024078 NP_067045 58497; 149428 HTCD37 4 4C NM_005506 NP_005497 950 SCARB2 5 4C NM_003713 NP_803133 8613 PPAP2B 5 4C NM_002247 NP_002987 6281 ST00A10 97 7 4C NM_002966 NP_005599 3043 HBB HBB HBB 1 4C NM_007081 NP_005744 22868 KIAA0971 98970 ST00A10 9889 HUMAUANT IN INTERCEPT 966 CD59 CD55	olgi autoantigen, golgin subfamily b. macropoloin		2804	NP_004478	NM_004487	4C	ncrc7131
8 4C NM_005013 NP_005004 4925; 399873 NUCB2 1 4C NM_017924 NP_060394 55017 C14orf119 3 4C NM_015411 NP_060394 55017 C14orf119 4C NM_025176 NP_0676983 79050 SUMF2 4C NM_024078 NP_076983 79050 MGC3162 4C NM_005506 NP_067045 58497; 149428 HTCD37 4C NM_003713 NP_803133 8613 PPAP2B 4C NM_002247 NP_002238 3778 KCNMA1 4C NM_002966 NP_002957 6281 S100A10 4C NM_002966 NP_005744 22868 KIAA0971 BB 4C NM_003333 NP_906076 966 CD59 CD59 4C NM_0033455 NP_003446 7753 ZNF202 ZNF202 4C NM_003418 NP_003409 51559 TU12B1-TY TU12B1-TY	protein phosphatase 1, regulatory subunit 7		5510	NP_002703	NM_002712	4C	1
4C NM_003013 NP_005004 4925; 39873 NUCB2 4C NM_017924 NP_060394 55017 C14orf119 4C NM_015411 NP_056226 25870 SUMF2 4C NM_005176 NP_076983 79050 MGC3162 4C NM_024078 NP_067045 58497; HTCD37 4C NM_003713 NP_803133 8613 PPAP2B 4C NM_003713 NP_00238 3778 KCNMA1 4C NM_002247 NP_002238 3778 KCNMA1 4C NM_002966 NP_002957 6281 S100A10 4C NM_003459 NP_005744 22868 KIAA0971 4C NM_007081 NP_009012 11158 RABL2B 4C NM_013285 NP_037417 29889 HUMAUANT 4C NM_016575 NP_033446 7753 ZNF202 Z 4C NM_006815 NP_036806 10959 TU12B1-TY T	zinc finger protein 9 (a cellular retroviral nucle		7555	NP_003409	NM_003418	4C	
4C NM_003013 NP_005004 4925; 399873 NUCB2 4C NM_017924 NP_060394 55017 C14orf119 4C NM_015411 NP_056226 25870 SUMF2 4C NM_005176 NP_076983 79050 MGC3162 4C NM_021222 NP_067045 58497; HTCD37 HTCD37 4C NM_003713 NP_803133 8613 PPAP2B 4C NM_002247 NP_002238 3778 KCNMA1 4C NM_000518 NP_002957 6281 S100A10 900A10 4C NM_000518 NP_0055744 22868 KIAA0971 11 4C NM_007081 NP_055744 22868 KIAA0971 11 4C NM_003285 NP_037417 29889 HUMAUANT IN ING 4C NM_013285 NP_037417 29889 HUMAUANT IN ING 4C NM_016575 NP_057659 51559 TU12B1-TY T	coated vesicle membrane protein		10959	NP_006806	NM_006815	4C	ncrc6981
4C NM_005013 NP_005004 4925; 39873 NUCB2 4C NM_017924 NP_060394 55017 C14orf119 4C NM_015411 NP_066226 25870 SUMF2 4C NM_025176 NP_076983 79050 MGC3162 4C NM_021222 NP_067045 58497; 58497; 149428 HTCD37 4C NM_003713 NP_803133 8613 PPAP2B 4C NM_002247 NP_002238 3778 KCNMA1 4C NM_002566 NP_002957 6281 S100A10 4C NM_007096 NP_005744 22868 KIAA0971 4C NM_007081 NP_005744 22868 KIAA0971 4C NM_007081 NP_009012 11158 RABL2B 4C NM_013285 NP_037417 29889 HUMAUANT n 4C NM_033455 NP_03446 7753 ZNF202 ZNF202			51559	NP_057659	NM_016575	4C	
8 4C NM_005013 NP_005004 4925; 39873 NUCB2 1 4C NM_017924 NP_060394 55017 C14orf119 2 4C NM_015411 NP_056226 25870 SUMF2 3 4C NM_005176 NP_076983 79050 MGC3162 4 C NM_024078 NP_076983 79050 MGC3162 4 C NM_024078 NP_076983 79050 MGC3162 4 C NM_0021222 NP_076983 79050 MGC3162 4 C NM_003713 NP_803133 8613 PPAP2B 4 C NM_002247 NP_002238 3778 KCNMA1 4 C NM_003518 NP_002957 6281 S100A10 S10A10 4 C NM_014929 NP_055744 22868 KIAA0971 HBB 4 C NM_007081 NP_090676 966 CD59 4 C NM_013285 NP_037417 29889 HUMAUANT n	zinc finger protein 202		7753	NP_003446	NM_003455	4C	
8 4C NM_005013 NP_005004 4925; 39873 NUCB2 1 4C NM_017924 NP_060394 55017 C14orf119 3 4C NM_015411 NP_056226 25870 SUMF2 4 C NM_024078 NP_056226 25870 SUMF2 4 C NM_024078 NP_076983 79050 MGC3162 5 4 C NM_021222 NP_067045 58497; 58497; 5950 HTCD37 4 C NM_003713 NP_803133 8613 PPAP2B 4 C NM_002247 NP_002238 3778 KCNMA1 4 C NM_002966 NP_002957 6281 S100A10 4 C NM_000518 NP_0055744 22868 KIAA0971 4 C NM_007081 NP_055744 22868 KIAA0971 4 C NM_007081 NP_06076 966 CD59	nucleolar GTPase	MAUANT	29889	NP_037417	NM_013285	46	
8 4C NM_005013 NP_005004 4925; 399873 NUCB2 1 4C NM_017924 NP_060394 55017 C14orf119 3 4C NM_015411 NP_056226 25870 SUMF2 4C NM_005176 NP_005167 517; 341412 ATP5G2 4C NM_024078 NP_076983 79050 MGC3162 4C NM_005506 NP_005497 950 SCARB2 4C NM_003713 NP_803133 8613 PPAP2B 4C NM_002247 NP_002238 3778 KCNMA1 4C NM_000518 NP_002957 6281 S100A10 4C NM_014929 NP_055744 22868 K1AA0971 4C NM_007081 NP_009012 11158 RABL2B	CD59 antigen p18-20 (antigen identified by monoclo		966	NP_976076	NM_203330	40	
8 4C NM_005013 NP_005004 4925; 399873 NUCB2 1 4C NM_017924 NP_060394 55017 C14orf119 3 4C NM_015411 NP_056226 25870 SUMF2 4C NM_005176 NP_076983 79050 MGC3162 3 4C NM_024078 NP_076983 79050 MGC3162 4C NM_025066 NP_067045 58497; HTCD37 HTCD37 4C NM_003713 NP_803133 8613 PPAP2B 4C NM_002247 NP_002238 3778 KCNMA1 4C NM_000518 NP_002957 6281 S100A10 4C NM_014929 NP_055744 22868 KIAA0971 PAMBB 4C NM_014929 NP_055744 22868 KIAA0971 PAMBB	RAB, member of RAS oncogene family-like 2B		111158	NP_009012	NM_00/081	40	- 1
4C NM_005013 NP_005004 4925; 39873 NUCB2 1 4C NM_017924 NP_060394 55017 C14orf119 3 4C NM_015411 NP_056226 25870 SUMF2 4 C NM_024078 NP_076983 79050 MGC3162 4 C NM_021222 NP_067045 58497; HTCD37 4 C NM_005506 NP_005497 950 SCARB2 4 C NM_003713 NP_803133 8613 PPAP2B 4 C NM_002966 NP_002957 6281 SCOARB2 4 C NM_000518 NP_000509 3043 HBB 4 C NM_014929 NP_055744 22868 KIAA0971	Similar to glutamate receptor, ionotropic, N-methy		339970		XM_291095	10 10	na=2617
4C NM_005013 NP_005004 4925; 399873 NUCB2 4C NM_017924 NP_060394 55017 C14orf119 3 4C NM_015411 NP_056226 25870 SUMF2 2 4C NM_005176 NP_056226 25870 SUMF2 3 4C NM_024078 NP_076983 79050 MGC3162 4C NM_021222 NP_067045 58497; HTCD37 HTCD37 4C NM_005506 NP_005497 950 SCARB2 4C NM_003713 NP_803133 8613 PPAP2B 4C NM_002966 NP_002957 6281 S100A10 4C NM_000518 NP_000509 3043 HBB	KIAA0971 protein	KIAA0971	22868	NP_055744	NM_014929	100	11C1C0302
8 4C NM_005013 NP_005004 4925; 399873 NUCB2 1 4C NM_017924 NP_060394 55017 C14orf119 3 4C NM_015411 NP_056226 25870 SUMF2 4C NM_005176 NP_005167 517; 341412 ATP5G2 4C NM_024078 NP_067045 58497; HTCD37 4C NM_005506 NP_067045 58497; HTCD37 4C NM_003713 NP_803133 8613 PPAP2B 4C NM_002966 NP_002957 6281 S100A10	hemoglobin, beta	НВВ	3043	NP_000509	81C000_IMN	à t	1/100101
4C NM_005013 NP_005004 4925; 39873 NUCB2 4C NM_017924 NP_060394 55017 C14orf119 4C NM_015411 NP_056226 25870 SUMF2 4C NM_024078 NP_076983 79050 MGC3162 4C NM_021222 NP_067045 58497; HTCD37 4C NM_005506 NP_005497 950 SCARB2 4C NM_003713 NP_803133 8613 PPAP2B 4C NM_002247 NP_002238 3778 KCNMA1	S100 calcium binding protein A10 (annexin II ligan	S100A10	6281	NP_002957	NM_002966	Å C	ncrc612/
4C NM_005013 NP_005004 4925; 399873 NUCB2 4C NM_017924 NP_060394 55017 C14orf119 4C NM_015411 NP_056226 25870 SUMF2 4C NM_005176 NP_005167 517; 341412 ATP5G2 4C NM_024078 NP_076983 79050 MGC3162 4C NM_021222 NP_067045 58497; HTCD37 4C NM_005506 NP_005497 950 SCARB2 4C NM_003713 NP_803133 8613 PPAP2B	potassium large conductance calcium-activated chan	KCNMAI	3778	NP_002238	NM_002247	40	ncrc6005
4C NM_005013 NP_005004 4925; 399873 NUCB2 4C NM_017924 NP_060394 55017 C14orf119 4C NM_015411 NP_056226 25870 SUMF2 4C NM_005176 NP_005167 517; 341412 ATP5G2 4C NM_024078 NP_076983 79050 MGC3162 4C NM_021222 NP_067045 58497; 149428 HTCD37 4C NM_005506 NP_005497 950 SCARB2	phosphatidic acid phosphatase type 2B	PPAP2B	8613	NP_803133	NM_003713	4C	ncrc5844
4C NM_005013 NP_005004 4925; 399873 NUCB2 4C NM_017924 NP_060394 55017 C14orf119 4C NM_015411 NP_056226 25870 SUMF2 4C NM_005176 NP_005167 517; 341412 ATP5G2 4C NM_024078 NP_076983 79050 MGC3162 4C NM_021222 NP_067045 58497; 58497; 149428 HTCD37	scavenger receptor class B, member 2	SCARB2	950	NP_005497	NM_005506	40	ncrc>806
NM_005013 NP_005004 4925; 399873 NUCB2 NM_017924 NP_060394 55017 C14orf119 NM_015411 NP_056226 25870 SUMF2 NM_005176 NP_005167 517; 341412 ATP5G2 NM_024078 NP_076983 79050 MGC3162	TcD37 homolog	HTCD37	58497; 149428	NP_067045	NM_021222	+	nc1c3/00
4C NM_005013 NP_005004 4925; 399873 NUCB2 4C NM_017924 NP_060394 55017 C14orf119 4C NM_015411 NP_056226 25870 SUMF2 4C NM_005176 NP_005167 517; 341412 ATP5G2	hypothetical protein MGC3162	MGC3162	79050	NP_076983	NM_024078	\$ 6	nc=c5760
4C NM_005013 NP_005004 4925; 399873 NUCB2 nucleobindin 4C NM_017924 NP_060394 55017 C14orf119 chromosome 4C NM_015411 NP_056226 25870 SUMF2 sulfatase moc	ATP synthase, H+ transporting, mitochondrial F0 co	f	517; 341412	NP_005167	NM_005176	1 6	mo=05750
4C NM_005013 NP_005004 4925; NUCB2 nucleobindin 4C NM_017924 NP_060394 55017 C14orf119 chromosome	sulfatase modifying factor 2	SUMF2	25870	NP_056226	NM_015411	† t	nereo663
NM_005013 NP_005004 4925; NUCB2 399873	chromosome 14 open reading frame 119	C14orf119	55017	NP_060394	NM_017924	t c	ncre5631
	nucleobindin 2	NUCB2	4925; 399873	NP_005004	NM_005013	4C	ncrc5608

FIP1 like 1 (S. cerevisiae)	FIPILI	8008	INI_1121/9			
Lloosomal protein L4			021C11 GIN	NM 030917	4C	ncrc9877 4
9	RPI 4	6124	NP_000959	NM_000968	4C	ncrc9867 4
chromosome I open reading frame 22	Clorf22	80267	NP_079467	NM_025191		
g Somore, 1 (FaiceIII)		23616			70	ncrc0055
lectin, galactoside-binding, soluble 1 (galactin	LGALS1 1	3956;	NP_002296	NM_002305	ŧ	
chromosome 1 open reading frame 24	Clorf24	1116496	NP_443198	NIX 002200	5 6	
v-maf musculoaponeurotic fibrosarcoma oncogene hom	MAF	4094	NP_00331	290C50 WIN	40	ncrc9758
secreted protein, acidic, cysteine-rich (osteonect	SPARC	6678	NP_003109	NM 005360	4C	ncrc9729
XTP3-transactivated protein B	ХТРЗТРВ	27248	JAE 020210	NIM 002110	4C	ncrc9727
basic transcription factor 3	BTF3	089; 220/93	NID 05/51/	NM 015701	4C	ncrc9712
		(80 2020	ND 001100	NM 001207	4C	ncrc9612
				NM_030778	4C	ncrc9557
17 - Transplant Const.				NM_025190	40	i
DEAD (Asp-Glu-Ala-Asp) box polynomide 24	DDX24	57062	NP_065147	NM_020414	÷	1
S100 calcium binding protein A4 (calcium protein	S100A4	6275	NP_062427	NM_002961	Š t	nor-0/102
splicing factor, arginine/serine-rich 5	SFRS5	6430	NP_008856	276000 IMM	70	ncrc0460 4C
hypothetical protein FLJ23221	FLJ23221	/9650	141 070000	SCOOL WIN	40	ncrc9428
nypothetical protein FLJ20035	1 LJZ0033	70000	ND 078855	NM 024579	4C	ncrc9368
	EI 12002 5	55601	NP 060101	NM_017631	4C	ncrc9331
CGL146 prot:	PNAS-4	51029	NP_057160	NM_016076	4C	ncrc9304
chromosome 9 open reading frame 121	C9orf121	158046	NP_660326	NM_145283	40	merey280
chromosome 20 open reading frame 64	C20orf64	112858	NP_291028	NM_033550	5	ncrc921/
1 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2				NM_024801	1 6	nere911/
chromosome 4 open reading frame 14	C4orf14	84273	NP_115689	NM_032313	† †C	ncrc9055
PR domain containing 10	PRDM10	56980	NP_955471	NM_020228		ncrc9023
development and differentiation	DDEF2	8853	NP_003878	NM_003887	4C	ncrc8976
CASP8 and EADD 1112	CFLAR	8837	NP_003870	NM_003879	4C	ncrc8903

proliferating cell nuclear antigen	PCNA	5111	NP_872590	NM_002592		500000
vacuolar protein sorting 29 (yeast)	VPS29	51699	NP 4/6528	091/CO_TAIN		sena0844 AC
myosin regulatory light chain MRLC2	MRLC2	103910	NP_291024	NIM 057180		se0a0799_4C
SRY (sex determining region Y)-box 4		0009	ND 201021	NW 033546	C	
Intestinal cell (MAK-like) kinase		6650	ND 002000	NM 003107		seoa0563 4C
11 11 12 12 12 12 12 12 12 12 12 12 12 1	ICK	22858	NP 057597	NM_016513	С	seoa0549 4C
BTG family member 3	BTG3	10950	NP_006797	NM_006806	4C	
sorting nexin 7	SNX7	51375	NP_689424	NM_015976	4°C	ı
small membrane protein 1	SMPI	23585	NP_055128	NM_014313	#	
laminin, gamma I (formerly LAMB2)	LAMCI	3915	NP_002284	NM 01223		\$20,000 20,000 20,000
transmembrane 4 superfamily member 6	TM4SF6	7105	NP_003261	NM 00202		sena0469 4C
myeloid cell nuclear differentiation antigen	MNDA	4332	NP_002423	NIM 002770		seoa0457 4C
vimentin	VIM	7431	NP_0033/1	NM 002422		seoa0288 4C
osteomodulin	OMD	4958	NP_005005	NM 003380		seoa0256 4C
thioredoxin	TXN	7295	NP_003320	MM_005017	10	sena0231 4C
reticulocalbin 2, EF-hand calcium binding domain	RCN2	5955	INP_002893	NIM 002200	40	seoa0221 4C
thymosin, beta 4, X-linked	I MSB4X	/114	THE 000000	COOCOO MIN	4C	seoa0219 4C
EGF-like-domain, multiple 6	E01.00	71114	ND 066022	NM 021109	4C	seoa0174 4C
LCD III III III III III III III III III I	EGEL &	25975	NP 056322	NM_015507	4C	seoa0172
RNA hinding motif protein 7	RBM7	10179	NP_057174	NM_016090	4C	seoa0111
succinate-CoA ligge ADD forming that it	SUCLA2	8803	NP_003841	NM_003850	4C	seoa0070
collagen, type I, alpha 2	COL1A2	1278	NP_000080	NM_000089	4C	
family with sequence similarity 3, member C	FAM3C	10447	NP_055703	NM_014888	40	
growth hormone inducible transmembrane protein	GHITM	27069	NP_055209	NM_014394	1 1 1 1	SC040008 4C
protease, serine, 11 (IGF binding)	PRSSII	5654	NP_002766	NM_002775	10 to	3C040003 4C
platelet-derived growth factor receptor, alpha pol	PDGFRA	5156	NP_006197	NM_006206	† C	G000000 4C
ALL1 fused gene from 5q31	AF5Q31	27125	NP_055238	NM_014423	4C	ncrc9899 4C

nucleotide binding protein 1 (MinD homolog, E. col	NUBPI	4682	NP_002475	NIVI_002484		750
matrilin 3	MATN3	4148 N	NP_002372	187 002181		Se0a 1700 Ar
H2A histone family, member Z	H2AFZ		INP_00209/	NM 002301		seoal 653 4C
mitochondrial ribosomal protein L13	MIKPLIS	°	NI OCCO	NM 002106		seoa1598 4C
Chromosomes 1-like	MDDI 12	28008	NP 054797	NM 014078	С	seoal 584 4C
SMC structural maintanance of 1	SMCILI	8243	NP_006297	NM_006306	C	seoa1582 4C
CD69 antigen (n60 early T-cell activation and	CD69	969	NP_001772	NM_001781	_ C	se0a13/1 4C
PRP4 pre-mRNA processing factor 4 homolog B (yeast	PRPF4B	8899	NP_789770	11 6000 TAIN		1
S100 calcium binding protein A4 (calcium protein,	S100A4		NP_062427	NIM 002013		seoal 559 4C
vesicle docking protein p115	VDP		NF_005/06	120000 MIN	C	seoal 552 4C
signal transducing adaptor molecule (SH3 domain an	STAM2	4	ND 002707	NM 003715	C	seoa1497 4C
tyrosine 3-monooxygenase/tryptophan 5-monooxygenas	YWHAZ	/534	NF_005/23	NM 005843	C	
chromosome 6 open reading frame 74	C6orf74		NF_05/14/	NM 145600	4C	
chromobox homolog 3 (HPI gamma homolog, Drosophila	CBX3		NP_05/6/1	NM 016063	4C	
ribosomal protein L5	RPL5	6123	NF 000900	NIM 016597	4C	seoa 173
mitochondrial ribosomal protein S33	MIKPOOS	01000	ND OOOCO	000000 NN	4C	seoal118
25 Pourentai pioteii MOC29891	MDDCCC	51650	NP 444263	NM 016071	†C	seoal 104 4C
hypothetical protein McCanona	MGC29891	126626	NP_653219	NM_144618	4C	seoal 102 4C
nascent-polypeptide-associated complex alpha alpha	NACA	4666; 5557	NP_005585	NM_005594	4C	seoa1089 4C
splicing factor, arginine/serine-rich 3	SFRS3	6428	NP_003008	NM_003017	4C	seoa1065
cyclin C	CCNC	892	NP_005181	061C00 WN	÷	
zinc finger RNA binding protein	ZFR	51663	NP_057191	NM_016107	j t	
p300/CBP-associated factor	PCAF	8850	NP_003875	NM_003884	<u>,</u>	Sec. 1036
calcyphosine	CAPS	828	NP_542157	NM_004058	å t	\$6000000
RAD23 homolog A (S. cerevisiae)	RAD23A	5886	NP_005044	NM_005053	A 6	se0a0931 4C
ribosomal protein L8	RPL8	6132	NP_150644	NM_033301	4C	seoa0926 4C
CD9 antigen (p24)	CD9	928	NP_001760	NM_001769	4C	seoa0913 4C

Secol 156 C							
9 4C NM_015907 NP_056991 51056 LAP3 6 4C NM_006292 NP_06692 TSG101 3 4C NM_006292 NP_060099 1738 DLD 0 4C NM_000108 NP_060023 7251 TSG101 2 4C NM_001219 NP_060152 113402 C6orf83 2 4C NM_018453 NP_060923 55837 C14orf11 4C NM_018453 NP_065189 57213 C13orf1 4C NM_020456 NP_065189 57213 C13orf1 4C NM_033014 NP_148935 4969 OGN 4C NM_033014 NP_996749 571; 54073 BACHI 4C NM_033014 NP_996749 571; 54073 BACHI 4C NM_0018491 NP_742000 150472; CBWD2 52 CALU NP_9872607 7324 UBE2EI 4C NM_001357 NP_085978 22920 KIFAP3 4C NM_006178 NP_085912 1	seoal /56	40	NM_002966	NP_002957	6281		
6 4C NM_006292 NP_006283 7251 TSG101 3 4C NM_000108 NP_000099 1738 DLD 3 4C NM_000108 NP_000099 1738 DLD 4C NM_001219 NP_060152 113402 C6orl83 9 4C NM_018453 NP_065090 23325 KIAA1033 1 4C NM_018453 NP_065189 57213 C13orf1 4C NM_020456 NP_965189 57213 C13orf1 4C NM_033014 NP_148935 4969 OGN 4C NM_020456 NP_996749 571: 54073 BACH1 4C NM_018491 NP_0148935 4969 OGN 4C NM_018491 NP_742000 150472: CBWD2 54C NM_003341 NP_872607 7324 UBE2E1 4C NM_003557 NP_005785 22920 KIFAP3 4C NM_006639 NP_00807 1660 DHX9 4C NM_006678 NP_0080630	seoal 789	4C	NM_015907	NP_056991	51056	LAP3	leucine aminopeptidase 3
3 4C NM_000108 NP_000099 1738 DLD 0 4C NM_145169 NP_660152 113402 C6orf83 0 4C NM_018459 NP_660152 113402 C6orf83 1 4C NM_018453 NP_060923 55837 C14orf11 1 4C NM_020456 NP_065189 57213 C13orf1 4 C NM_033014 NP_148935 4969 OGN 4 C NM_020866 NP_996749 571; 54073 BACH1 4 C NM_018491 NP_742000 150472; CBWD2 5 4C NM_018491 NP_872607 7324 UBE2E1 4 C NM_002791 NP_872607 7324 UBE2E1 4 C NM_003341 NP_985785 22920 KIFAP3 4 C NM_0014970 NP_085077 1660 DHX9 4 C NM_007281 NP_008507 1660 DHX9 4 C NM_006639 NP_003606 9497 SLC4A7 4 C NM_006678 NP_0036169	seoa1856	4C	NM_006292	NP_006283	7251		tumor susceptibility gene 101
0 4C NM_145169 NP_660152 113402 C6orf83 9 4C NM_001219 NP_0601210 813 CALU 9 4C NM_018453 NP_060923 55837 C14orf11 1 4C NM_018275 NP_065990 23325 KIAA1033 2 4C NM_020456 NP_065189 57213 C13orf1 4 C NM_033014 NP_148935 4969 OGN 4 C NM_033014 NP_996749 571; 54073 BACH1 4 C NM_018491 NP_996749 571; 54073 BACH1 4 C NM_018491 NP_901016 6228 RPS23 4 C NM_018491 NP_872607 7324 UBE2E1 4 C NM_003341 NP_872607 7324 UBE2E1 4 C NM_001357 NP_085077 1660 DHX9 4 C NM_006639 NP_0055785 22920 KIFAP3 4 C NM_006639 NP_003606 9497 SLC4A7 4 C NM_006639		4C	NM_000108	NP_000099	1738		dihydrolipoamide dehydrogenase (E3 component of py
9 4C NM_001219 NP_001210 813 CALU 2 4C NM_018453 NP_060923 55837 C14orf11 1 4C NM_018453 NP_060923 55837 C14orf11 2 4C NM_020456 NP_065189 57213 C13orf1 4 C NM_033014 NP_148935 4969 OGN 4 C NM_033014 NP_996749 571; 54073 BACH1 4 C NM_018491 NP_001016 6228 RPS23 4 C NM_018491 NP_742000 150472; CBWD2 5 4C NM_003341 NP_872607 7324 UBE2E1 4 C NM_002791 NP_085077 1660 DHX9 4 C NM_006639 NP_085077 1660 DHX9 4 C NM_007281 NP_008630 10800 CYSLTRI 4 C NM_003615 NP_003606 9497 SLC4A7 4 C NM_006737 NP_0036169 4905 NSF 4 C NM_006827 NP_005	seoa1940	4C	NM_145169	NP_660152	113402		chromosome 6 open reading frame 83
2 4C NM_018453 NP_060923 55837 C14orf11 1 4C NM_015275 NP_056090 23325 KIAA1033 2 4C NM_020456 NP_055189 57213 C13orf1 4 C NM_033014 NP_148935 4969 OGN 4 C NM_018491 NP_996749 571; 54073 BACH1 4 C NM_018491 NP_001016 6228 RPS23 4 C NM_018491 NP_742000 150472; CBWD2 55871 S5871 CBWD2 4 C NM_003341 NP_872607 7324 UBE2E1 4 C NM_00357 NP_085785 22920 KIFAP3 4 C NM_006639 NP_085077 1660 DHX9 4 C NM_003615 NP_085077 1660 DHX9 4 C NM_003615 NP_003606 9497 SLC4A7 4 C NM_003545 NP_031401 23438; DISCA7 TARDBP 4 C NM_006827 NP_006818 10972 TMP21	seoa1979	4C	NM_001219	NP_001210	813		calumenin
1 4C NM_015275 NP_056090 23325 KIAA1033 2 4C NM_020456 NP_065189 57213 C13orf1 5 4C NM_020456 NP_065189 57213 C13orf1 4 C NM_033014 NP_148935 4969 OGN 4 C NM_01025 NP_091016 6228 RPS23 4 C NM_018491 NP_742000 150472; CBWD2 5 871 S5871 UBE2E1 4 C NM_003341 NP_872607 7324 UBE2E1 4 C NM_014970 NP_055785 22920 KIFAP3 4 C NM_00639 NP_085077 1660 DHX9 4 C NM_006639 NP_0955785 22920 KIFAP3 4 C NM_006639 NP_008507 1660 DHX9 4 C NM_003615 NP_003606 9497 SLC4A7 4 C NM_006178 NP_0031401 23435; TARDBP 4 C NM_005927 NP_006818 10972 TMP21 <td>seoa2012</td> <td>4C</td> <td>NM_018453</td> <td>NP_060923</td> <td>55837</td> <td></td> <td></td>	seoa2012	4C	NM_018453	NP_060923	55837		
2 4C NM_020456 NP_065189 57213 C13orf1 5 4C NM_033014 NP_148935 4969 OGN 1 4C NM_033014 NP_996749 571; 54073 BACH1 4 C NM_01025 NP_001016 6228 RPS23 4 C NM_018491 NP_742000 150472; CBWD2 CBWD2 55871 NP_002782 5687 PSMA6 4 C NM_003341 NP_085077 1660 DHX9 4 C NM_0014970 NP_085077 1660 DHX9 4 C NM_006639 NP_085077 1660 DHX9 4 C NM_003615 NP_0036630 10800 CYSLTRI 4 C NM_003615 NP_003606 9497 SLC4A7 4 C NM_006178 NP_006169 4905 NSF 4 C NM_006827 NP_006818 10972 TMP21 4 C NM_006827 NP_006818 10972 TMP21	seoa2041	4C	NM_015275	NP_056090	23325		
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4C NM_206866 NP_996749 571; 54073 BACHI 2 4C NM_001025 NP_001016 6228 RPS23 2 4C NM_018491 NP_742000 150472; CBWD2 CBWD2 3 4C NM_003341 NP_872607 7324 UBE2E1 4C NM_002791 NP_05785 22920 KIFAP3 4C NM_014970 NP_085077 1660 DHX9 4C NM_001357 NP_085077 1660 DHX9 4C NM_006639 NP_0036630 10800 CYSLTRI 4C NM_003615 NP_003666 9497 SLC4A7 4C NM_005545 NP_0036169 4905 NSF 4C NM_006827 NP_005918 23435; TARDBP 4C NM_006827 NP_006818 10972 TMP21		4C	NM_033014				osteoglycin (osteoinductive factor, mimecan)
4C NM_001025 NP_001016 6228 RPS23 4C NM_018491 NP_742000 150472; CBWD2 5871 55871 CBWD2 4C NM_003341 NP_872607 7324 UBE2E1 4C NM_002791 NP_005785 22920 KIFAP3 4C NM_014970 NP_085077 1660 DHX9 4C NM_006639 NP_085077 1660 DHX9 4C NM_007281 NP_009212 11341 SCRG1 4C NM_003615 NP_003606 9497 SLC4A7 4C NM_006178 NP_006169 4905 NSF 4C NM_007375 NP_031401 23435; TARDBP 4C NM_006827 NP_006818 10972 TMP21 4C NM_006827 NP_006818 10972 TMP21 1		4C	NM_206866	NP_996749			3TB and CNC homology 1, basic leucine zipper trans
2 4C NM_018491 NP_742000 150472; 55871 CBWD2 4 C NM_003341 NP_872607 7324 UBE2E1 4 C NM_002791 NP_002782 5687 PSMA6 4 C NM_014970 NP_085077 1660 DHX9 4 C NM_001357 NP_085077 1660 DHX9 4 C NM_006639 NP_009212 11341 SCRGI 4 C NM_003615 NP_003606 9497 SLC4A7 4 C NM_005545 NP_958934 3671 ISLR 4 C NM_006178 NP_006169 4905 NSF 4 C NM_007375 NP_031401 23435; TARDBP 4 C NM_005927 NP_006818 10972 TMP21 1		4C	NM_001025				ibosomal protein S23
4C NM_003341 NP_872607 7324 UBE2E1 4C NM_002791 NP_002782 5687 PSMA6 4C NM_014970 NP_055785 22920 KIFAP3 4C NM_001357 NP_085077 1660 DHX9 4C NM_006639 NP_009212 11341 SCRG1 4C NM_007281 NP_009212 11341 SCRG1 4C NM_003615 NP_003606 9497 SLC4A7 4C NM_006178 NP_006169 4905 NSF 4C NM_007375 NP_031401 23435; TARDBP 4C NM_005927 NP_006818 10972 TMP21 4C NM_006827 NP_006818 10972 TMP21			NM_018491		2;		COBW domain containing 2
4C NM_002791 NP_002782 5687 PSMA6 4C NM_014970 NP_055785 22920 KIFAP3 4C NM_001357 NP_085077 1660 DHX9 4C NM_006639 NP_006630 10800 CYSLTRI 4C NM_007281 NP_003606 9497 SLC4A7 4C NM_005545 NP_958934 3671 ISLR 4C NM_006178 NP_006169 4905 NSF 4C NM_007375 NP_031401 23435; TARDBP 4C NM_005927 NP_006818 10972 TMP21 4C NM_006827 NP_006818 10972 TMP21			NM_003341	NP_872607			biquitin-conjugating enzyme E2E 1 (UBC4/5 homolog
4C NM_014970 NP_055785 22920 KIFAP3 4C NM_001357 NP_085077 1660 DHX9 4C NM_006639 NP_006630 10800 CYSLTRI 4C NM_007281 NP_009212 11341 SCRG1 4C NM_003615 NP_003606 9497 SLC4A7 4C NM_006178 NP_958934 3671 ISLR 4C NM_007375 NP_006169 4905 NSF 4C NM_005927 NP_005918 4238 MFAP3 4C NM_006827 NP_006818 10972 TMP21	seoa2381		NM_002791				roteasome (prosome, macropain) subunit, alpha typ
4C NM_001357 NP_085077 1660 DHX9 4C NM_006639 NP_006630 10800 CYSLTR1 4C NM_007281 NP_009212 11341 SCRG1 4C NM_003615 NP_003606 9497 SLC4A7 4C NM_005545 NP_958934 3671 ISLR 4C NM_006178 NP_006169 4905 NSF 4C NM_007375 NP_031401 23435; TARDBP 4C NM_005927 NP_005918 4238 MFAP3 4C NM_006827 NP_006818 10972 TMP21	seoa2391		NM_014970				inesin-associated protein 3
4C NM_00639 NP_00630 10800 CYSLTRI 4C NM_007281 NP_009212 11341 SCRG1 4C NM_003615 NP_003606 9497 SLC4A7 4C NM_005545 NP_958934 3671 ISLR 4C NM_006178 NP_006169 4905 NSF 4C NM_007375 NP_031401 23435; TARDBP 4C NM_005927 NP_005918 4238 MFAP3 4C NM_006827 NP_006818 10972 TMP21	3		NM_001357	NP_085077			DEAH (Asp-Glu-Ala-His) box polypeptide 9
4C NM_007281 NP_009212 11341 SCRGI 4C NM_003615 NP_003606 9497 SLC4A7 4C NM_005545 NP_958934 3671 ISLR 4C NM_006178 NP_006169 4905 NSF 4C NM_007375 NP_031401 23435; 391494 TARDBP 4C NM_005927 NP_005918 4238 MFAP3 4C NM_006827 NP_006818 10972 TMP21	I		NM_006639				ysteinyl leukotriene receptor l
4C NM_003615 NP_003606 9497 SLC4A7 4C NM_005545 NP_958934 3671 ISLR 4C NM_006178 NP_006169 4905 NSF 4C NM_007375 NP_031401 23435; 391494 TARDBP 4C NM_005927 NP_005918 4238 MFAP3 4C NM_006827 NP_006818 10972 TMP21	L		NM_007281				crapie responsive protein 1
4C NM_005545 NP_958934 3671 ISLR 4C NM_006178 NP_006169 4905 NSF 4C NM_007375 NP_031401 23435; TARDBP 4C NM_005927 NP_005918 4238 MFAP3 4C NM_006827 NP_006818 10972 TMP21			NM_003615				olute carrier family 4, sodium bicarbonate cotran
4C NM_006178 NP_006169 4905 NSF 4C NM_007375 NP_031401 23435; TARDBP 4C NM_005927 NP_005918 4238 MFAP3 4C NM_006827 NP_006818 10972 TMP21	seoa2639 4						nmunoglobulin superfamily containing leucine-rich
4C NM_007375 NP_031401 23435; TARDBP TAR DNA bind 4C NM_005927 NP_005918 4238 MFAP3 microfibrillar-a 4C NM_006827 NP_006818 10972 TMP21 transmembrane	seoa2641 4						ethylmaleimide-sensitive factor
4C NM_005927 NP_005918 4238 MFAP3 microfibrillar-a 4C NM_006827 NP_006818 10972 TMP21 transmembrane	1						AR DNA binding protein
4C NM_006827 NP_006818 10972 TMP21 transmembrane	1						nicrofibrillar-associated protein 3
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1) or appropriate the state of	3	シンメ	- CC4XD2	1,040,7		
cytochrome c oxidase subunit VIIa polypentide 2 (1	COX7A2		NE_001020	NM 004872	4C	
ubiquitin B	dao	13.41	NID 001856	NM 001865	4C	seoa4708 4
ruoumi, aipiid, ubiquitous	I IBB	7314	NP 061828	NM_018955	4C	seoa4681 '
This is also the second service and the secon	K-ALPHA-I	10376	NP_006073	NM_006082	C	seoa4600 4C
RABIA, member RAS oncovene family	RABIA	5861	NP_004152	NM_004161	4C	se0a4586
CGI-128 protein	CGI-128	51647	NP_057146	NM_016062	; *	
choroideremia (Rab escort protein 1)	СНМ	11121	NP_000381	NM_000390	5 7	
potassium voltage-gated channel, delayed-rectifier	KCNS3	3790	NP_002243	NM_00222	7	Se094518
cell cycle progression 8 protein	CPR8	9236	NP_065790	NM_020739		-
peptidase D	PEPD	5184	NP_000276	NM_000285		SCO4+452 4C
ubiquinol-cytochrome c reductase hinge protein	UQCRH	7388	NP_005995	NM_006004	6	seoa44593 4C
tuberous sclerosis 2	TSC2	7249	NP_066400	NM_000548	4	3004300
collagen, type III, alpha 1 (Ehlers-Danlos syndrom	COL3A1	1281	NP_000081	NM_000090	40	
nucleolar protein 5A (56kDa with KKE/D repeat)	NOL5A	10528	NP_006383	NM_006392	+	
hypothetical protein FLJ20507	FLJ20507	55654	NP_060319	NW 01/649	5	saca/26/
ceroid-lipofuscinosis, neuronal 2, late infantile	CLN2	1200	NP_000382	NW 017840	4C	seoa4167
actin related protein 2/3 complex, subunit 3, 21kD	ARPC3	10094	NP_005710	NIM 000301	40	seoa4158 4C
clathrin, light polypeptide (Lca)	CLTA	1211	NP_009027	NM 005710	40 6	seoa4107
CD74 antigen (invariant polypeptide of major histo	CD74	972	NP_004346	NM_004355	7 7	se0a4040
development and differentiation enhancing factor 2	DDEF2	8853	NP_003878	NM_003887	7	Seca4040
retinoic acid induced 2	RAI2	10742	NP_068557	28/170 MM	5 8	sena/1022
oxytocin receptor	OXTR	5021	NP_00090/	MM 001765	40	se0a4017
eukaryotic translation initiation factor 4A, isofo	EIF4A2	1974	NP_001958	NW 000016	4C 8	seoa3910 4C
retinol dehydrogenase 11 (all-trans and 9-cis)	RDH11	51109; 26101	NP_057110	NM_016026	40	seoa3847 4C
voltage-dependent anion channel 2	VDAC2	7417	NP_003366	NM_003375	6	Se0a3/1/ 4C

myosin IB	MY01B	4430	NP_036355	NM_012223	4C	seoa6197
farnesyltransferase, CAAX box, alpha	FNTA	2339	NP_002018	NM_002027	4C	
ras homolog gene family, member B	RHOB	388 .	NP_004031	NM_004040	4C	seoa6152
tumor protein p53 (Li-Fraumeni syndrome)	TP53	7157	NP_000537	NM_000546	4C	seoa6137
progesterone receptor membrane component 1	PGRMCI	10857	NP_006658	NM_006667	4C	
zinc finger protein 146	ZNF146	7705	NP_009076	NM_007145	4C	seoa6038
hypothetical protein MGC10067	MGC10067	134510	NP_659486	NM_145049	4C	seoa6032
CAP, adenylate cyclase-associated protein 1 (yeast	CAPI	10487	NP_006358	NM_006367	4C	seoa5986 4C
OGT(O-Glc-NAc transferase)-interacting protein 106	OIP106	22906	NP_055780	NM_014965	4C	seoa5933
ornithine decarboxylase antizyme 1	OAZI	4946	NP_004143	NM_004152	40	seoa5911
SON DNA binding protein	SON	6651	NP_620305	NM_058183	4C	seoa5894 4C
axin 2 (conductin, axil)	AXIN2	8313	NP_004646	NM_004655	4C	seoa5742
methionyl aminopeptidase 2	METAP2	10988	NP_006829	NM_006838	4C	seoa5683 4C
platelet-activating factor acetylhydrolase, isofor	PAFAHIBI	5048	NP_000421	NM_000430	4C	seoa5662
strand-exchange protein 1	1-Sep	54464	NP_061874	NM_019001	4C	
eukaryotic translation initiation factor 3, subuni	EIF3S3	8667	NP_003747	NM_003756	4C	seoa5577 4C
leukotriene A4 hydrolase	LTA4H	4048	NP_000886	NM_000895	4C	seoa5554 4C
NADH dehydrogenase (ubiquinone) Fe-S protein 5, 15	NDUFS5	4725	NP_004543	NM_004552	4C	seoa5547 4C
F-box and leucine-rich repeat protein 3A	FBXL3A	26224	NP_036290	NM_012158	4C	seoa5465
heat shock 70kDa protein 5 (glucose-regulated prot	HSPA5	3309	NP_005338	NM_005347	4C	seoa5429
oxysterol binding protein-like 1A	OSBPLIA	114876	NP_579802	NM_080597	4C	seoa5382
ribosomal protein L17	RPL17	6139	NP_000976	NM_000985	4C	seoa5302
nucleoporin 54kDa	NUP54	53371	NP_059122	NM_017426	4C	seoa5235
synaptophysin-like protein	SYPL	6856	NP_874384	NM_006754	4C	seoa5090
comparative gene identification transcript 37	CGI-37	51388	NP_057185	NM_016101	4C	seoa4727 4C
					T	

guanine nucleotide binding protein (G protein), ga	GNG11 g	2791	NP_004117	NM_004126		3504/4/6 40
period homolog 2 (Drosophila)	PER2 p	8864	NP_073728	NM_022817		seoa/4/4 4C
core-binding factor, beta subunit	CBFB c	865	NP_074036	NM_001755	4C	seoa7459 "
sorting nexin 6	s 9XNS	58533	NP_689419	NM_021249	4C	
proteasome (prosome, macropain) activator subunit	PSME1 p	5720	NP_788955	NM_006263	4C	
hypothetical protein FLJ36812	FLJ36812 h	255252	NP_694992	NM_153260	40	
proline-serine-threonine phosphatase interacting p	PSTPIP2 F	9050	NP_077748	NM_024430		
HT001 protein	HT001	28990	NP_054784	NM_014065	4C	seoa7223
FK506 binding protein 9, 63 kDa	FKBP9	11328	NP_009201	NM_007270	†C	seoa7212 4C
vinculin	VCL	7414	NP_054706	NM_014000	4C	seoa7178 4C
ring finger protein 141	RNF141	50862	NP_057506	NM_016422	4C	seoa7094 4C
nucleoredoxin	NXN	64359	NP_071908	NM_022463	4C	seoa6923
low molecular mass ubiquinone-binding protein (9.5	QP-C	27089	NP_055217	NM_014402	4C	1
testis enhanced gene transcript (BAX inhibitor 1)	TEGT	7009	NP_003208	NM_003217	4C	
a disintegrin-like and metalloprotease (reprolysin	ADAMTSI	9510	NP_008919	NM_006988	4C	
checkpoint with forkhead and ring finger domains	CHFR	55743	NP_060693	NM_018223	4C	
matrix metalloproteinase 11 (stromelysin 3)	MMP11	4320	NP_005931	NM_005940	4C	seoa6621
coronin, actin binding protein, 1C	COROIC	23603	NP_055140	NM_014325	40	
apoptosis inhibitor 5	API5	8539	NP_006586	NM_006595	4C	
SET and MYND domain containing 3	SMYD3	64754	NP_073580	NM_022743	4C	
sorting nexin 3	SNX3	8724	NP_690041	NM_152828	4C	1
UMP-CMP kinase	UMP-CMPK	51727	NP_057392	NM_016308	4C	1
inhibitor of growth family, member 1	INGI	3621	NP_937862	NM_198219	4C	se0a6393
synaptotagmin binding, cytoplasmic RNA interacting	SYNCRIP	10492; 57231	NP_006363	NM_153816	4C	
ubiquitin specific protease 33	USP33	23032	NP_963920	NM_015017	40	Se080313 4C
WWW				101	70	2167000

HIV-1 1at interactive protein 2, 30kDa	HIAIIP2 HIV	10333 Н17	NF_000401	NIVI_000410	+	35048040
SPARC related modular calcium binding 1	3				4C	
13kDa differentiation-associated protein		55967 DAP13			4C	1
ubiquitin specific protease 9, X-linked (fat facet		8239 USP9X	NP_068706 8:	NM_004652	4C	seoa8556
NADH dehydrogenase (ubiquinone) Fe-S protein 4, 18	NDUFS4 NAI	4724 NDI	NP_002486 4	NM_002495	4C	seoa8543
small nuclear ribonucleoprotein polypeptide G		6637 SNRPG	NP_003087 60	NM_003096	4C	seoa8486
type I transmembrane C-type lectin receptor DCL-1		9936 DCL-1	NP_055695 9	NM_014880	4C	seoa8443
t-complex-associated-testis-expressed 1-like 1	TCTEL1 t-cor	TCT (6993	NP_006510 6	NM_006519	4C	seoa8401 4C
hydroxysteroid (17-beta) dehydrogenase 4	HSD17B4 hydr	3295 HSI	NP_000405 3:	NM_000414	4C	seoa8399
progesterone receptor membrane component 2	PGRMC2 prog	10424 PGF	NP_006311 10	NM_006320	4C	seoa8397
ornithine aminotransferase (gyrate atrophy)		4942 OAT	NP_000265 4	NM_000274	4C	seoa8348
proteasome (prosome, macropain) subunit, alpha typ		5685 PSMA4	NP_002780 5	NM_002789	4C	seoa8300
activity-dependent neuroprotector		23394 ADNP	NP_852107 2:	NM_015339	4C	seoa8239
ATP synthase, H+ transporting, mitochondrial F0 co		522 ATP5J	NP_001676 5:	NM_001685	4C	seoa8232
catenin (cadherin-associated protein), alpha 1, 10	CTNNAI cater	1495; CTN 401210	NP_001894 1.	NM_001903	4C	seoa8177
Norrie disease (pseudoglioma)		4693 NDP	NP_000257 4	NM_000266	4C	seoa7935
cisplatin resistance-associated overexpressed prot	LUC7A cisp	51747; LUC 10414	NP_057508 5	NM_016424	4C	seoa7902
jumping translocation breakpoint		10899 ЈТВ	NP_006685	NM_006694	4C	seoa7626
putative breast adenocarcinoma marker (32kD)		27243 BC-2	NP_940818 2	NM_014453	4C	seoa7608
TANK-binding kinase 1		29110 TBK1	NP_037386 2	NM_013254	4C	seoa7587
				NM_003138	4C	seoa7555
likely ortholog of mouse monocyte macrophage 19	MMRP19 likel	51074 MN	NP_057041 5	NM_015957	4C	seoa7546 4C
lin	translin	7247 TSN	NP_004613 7	NM_004622	4C	seoa7530 4C
Notch homolog 2 (Drosophila)	NOTCH2 Note	4853 NO	NP_077719 4	NM_024408	4C	seoa7517
RRS1 ribosome biogenesis regulator homolog (S. cer		23212 RRS1	NP_055984 2	NM_015169	4C	seoa7509

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GABA(A) receptor-associated protein	GABARAP (11337; 79142; 23399;	NP_009209	NM_015343	4C	seoa9814 4C
CGG triplet repeat binding protein 1	CGGBP1 (8545; 401078	NP_003654	NM_003663	† †C	seoa9740 4C
epimorphin	EPIM	2054	NP_919337	NM_194356	4C	seoa9739 4C
chromobox homolog 3 (HP1 gamma homolog, Drosophila	CBX3	11335	NP_057671	NM_016587	4C	seoa9729 4C
mitochondrial ribosomal protein S30	MRPS30	10884	NP_057724	NM_016640	4C	seoa9724 4C
adenylate kinase 3 like 1	AK3L1	50808	NP_057366	NM_016282	4C	seoa9711 4C
tumor necrosis factor (ligand) superfamily, member	TNFSF11	8600	NP_143026	NM_003701	4C	seoa9619 4C
senescence downregulated leo1-like	LOC123169	123169	NP_620147	NM_138792	4C	seoa9537
piwi-like 2 (Drosophila)	HIWI2	143689	NP_689644	NM_152431	4C	seoa9482
chromosome 6 open reading frame 166	C60rf166	55122	NP_060534	NM_018064	4C	seoa9389
lysosomal-associated membrane protein 2	LAMP2	3920	NP_054701	NM_013995	4C	seoa9373
growth and transformation-dependent protein	E2IG5	26355	NP_055182	NM_014367	4C	seoa9357
amyotrophic lateral sclerosis 2 (juvenile) chromos	ALS2CR3	66008	NP_055864	NM_015049	4C	1
unactive progesterone receptor, 23 kD	TEBP	10728	NP_006592	NM_006601	4C	seoa9131 4C
zinc finger protein 302	ZNF302	55900	NP_061145	NM_018675	4C	seoa9042 4C
chromosome 2 open reading frame 3	C2orf3	6936	NP_003194	NM_003203	4C	seoa8979 4C
GNAS complex locus	GNAS	2778	NP_536351	NM_080425	4C	seoa8916 4C
cytochrome c, somatic	CYCS	54205	NP_061820	NM_018947	4C	seoa8912
heat shock factor binding protein 1	HSBP1	3281	NP_001528	NM_001537	4C	seoa8902
				NM_002450	4C	seoa8894
ankyrin repeat, family A (RFXANK-like), 2	ANKRA2	57763	NP_075526	NM_023039	4C	seoa8870 4C
ribosomal protein S27a	RPS27A	6233	NP_002945	NM_002954	4C	seoa8814
glioma tumor suppressor candidate region gene 2	GLTSCR2	29997	NP_056525	NM_015710	4C	seoa8754 4C
steroid sensitive gene 1	URB	151887	NP_955806	NM_199511	4C	seoa8642

4C NM_001424 NP_001415 2013 EMP2 epithelial me 4C NM_003758 NP_003749 8669 EIF3S1 eukaryotic tr 4C NM_003758 NP_078813 79587 FLJ12118 hypothetical 4C NM_004894 NP_063792 5556 C14ort2 chromosome 4C NM_004894 NP_060792 55776 C6ort64 chromosome 4C NM_002106 NP_060792 55776 C6ort64 chromosome 4C NM_005034 NP_060968 9988: DMTF1 cyclin D bind 4C NM_005034 NP_066968 9988: DMTF1 cyclin D bind 4C NM_004553 NP_004544 4726 NDUFS6 NADH dehyd 4C NM_004572 NP_0055787 22980 KIAA1049 KIAA1049 pt 4C NM_004501 NP_006592 10728 TEBP unactive prog 4C NM_007097 NP_006592 10728 HFI H factor I (cc	myouopiiii	17		111 -00000			
4C NM_001424 NP_001415 2013 EMP2 4C NM_003758 NP_003749 8669 EIF3SI 4C NM_024537 NP_078813 79587 FLJ12118 4C NM_04894 NP_004885 9556 C14orf2 4C NM_018322 NP_060792 55776 C6orf64 4C NM_005034 NP_066968 9988; DMTF1 4C NM_004553 NP_065968 9988; DMTF1 4C NM_004553 NP_004544 4726 NDUFS6 4C NM_004553 NP_0050969 9349 RPL23 4C NM_006601 NP_006592 10728 TEBP 4C NM_001497 NP_006996 9349 RPL23 4C NM_001499 NP_00177 307	1 7	MTDM	310	ND 665807	NM 145808	5	
4C NM_001424 NP_001415 2013 EMP2 5 4C NM_003758 NP_003749 8669 EIF3SI 6 4C NM_024537 NP_078813 79587 FLJ12118 6 4C NM_004894 NP_060792 8409 UXT 6 4C NM_004894 NP_060792 55776 C6orf64 6 4C NM_002106 NP_060792 55776 C6orf64 7 4C NM_002104 NP_060968 9988; DMTF1 7 4C NM_0030938 NP_112200 81671 VMP1 7 4C NM_004523 NP_004544 4726 NDUFS6 7 4C NM_004572 NP_0055787 22980 KIAA1049 7 4C NM_006601 NP_006592 10728 TEBP 7 4C NM_006601 NP_006592 10728 TEBP 7 4C NM_0014971 NP_006592 10728 TEBP 7 4C NM_0014991 NP_006969 301 ACTN1 7 4C NM_004454 <td< td=""><td>cell division cycle 42 (CTB Linding and in 25)</td><td>CDC42</td><td>998</td><td>NP 426359</td><td>NM 001791</td><td>Toll</td><td>seob0263 4</td></td<>	cell division cycle 42 (CTB Linding and in 25)	CDC42	998	NP 426359	NM 001791	Toll	seob0263 4
4C NM_001424 NP_001415 2013 EMP2 4C NM_003758 NP_003749 8669 EIF3S1 4C NM_024537 NP_078813 79587 FLJ12118 4C NM_0183477 NP_070582 8409 UXT 4C NM_018322 NP_060792 55776 C6orf64 4C NM_002106 NP_002097 3015 H2AFZ 4C NM_005034 NP_066968 9988; DMTF1 4C NM_030938 NP_112200 81671 VMP1 4C NM_004553 NP_0055787 22980 KIAA1049 4C NM_004553 NP_0055787 22980 KIAA1049 4C NM_006601 NP_006592 10728 TEBP 4C NM_0014972 NP_006592 10728 TEBP 4C NM_0006601 NP_006592 10728 TEBP 4C NM_0014991 NP_006993 87 ACTN1 4C NM_005005 NP_004996 <td< td=""><td>ets variant gene 5 (ets-related molecule)</td><td>ETV5</td><td>2119</td><td>NP_004445</td><td>NM_004454</td><td>Ť.</td><td>seob0253 </td></td<>	ets variant gene 5 (ets-related molecule)	ETV5	2119	NP_004445	NM_004454	Ť.	seob0253
4C NM_001424 NP_001415 2013 EMP2 4C NM_003758 NP_003749 8669 EIF3S1 4C NM_024337 NP_078813 79587 FLJ12118 4C NM_0183477 NP_070582 8409 UXT 4C NM_018322 NP_060792 5576 C14ort2 4C NM_002106 NP_002097 3015 H2AFZ 4C NM_005034 NP_005025 5440 POLR2K 4C NM_005034 NP_066968 9988; 402563; 401385 DMTF1 4C NM_030938 NP_112200 81671 VMP1 4C NM_004553 NP_004544 4726 NDUFS6 4C NM_004572 NP_0055787 22980 KIAA1049 4C NM_000978 NP_006592 10728 TEBP 4C NM_000102 NP_006592 10728 TEBP 4C NM_001102 NP_0001093 87 ACTN1 4C NM_014991 NP_848700	NADH dehydrogenase (ubiquinone) 1 beta subcomplex,	NDUFB9	4715	NP_004996	NM_005005	ţĊ	seob0248
4C NM_001424 NP_001415 2013 EMP2 4C NM_003758 NP_003749 8669 EIF3S1 4C NM_024537 NP_078813 79587 FLJ12118 4C NM_004894 NP_004885 9556 C14orf2 4C NM_018322 NP_060792 55776 C6orf64 4C NM_002106 NP_002097 3015 H2AFZ 4C NM_005034 NP_005025 5440 POLR2K 4C NM_005034 NP_066968 9988: 401385 DMTF1 4C NM_030938 NP_112200 81671 VMP1 4C NM_004553 NP_002183 3624 INHBA 4C NM_004553 NP_004544 4726 NDUFS6 4C NM_006601 NP_006592 10728 TEBP 4C NM_000102 NP_0001093 87 ACTN1 4C NM_000102 NP_001093 87 ACTN1	WD repeat and FYVE domain containing 3	WDFY3	23001	NP_848700	NM_014991	†C	seob0228
NM_001424 NP_001415 2013 EMP2 NM_003758 NP_003749 8669 EIF3S1 NM_0024537 NP_078813 79587 FLJ12118 NM_004894 NP_004885 9556 C14orf2 NM_002106 NP_0060792 55776 C6orf64 NM_002106 NP_005025 5440 POLR2K NM_021145 NP_066968 9988; DMTF1 NM_030938 NP_112200 81671 VMP1 NM_014972 NP_004544 4726 NDUFS6 NM_006601 NP_00699 9349 RPL23 NM_000186 NP_000691 301 ANXA1	actinin, alpha 1	ACTN1	87	NP_001093	NM_001102	†C	seob0201
4C NM_001424 NP_001415 2013 EMP2 4C NM_003758 NP_003749 8669 EIF3S1 4C NM_0024537 NP_078813 79587 FLJ12118 4C NM_004894 NP_004885 9556 C14orf2 4C NM_004894 NP_060792 55776 C6orf64 4C NM_002106 NP_060792 55776 C6orf64 4C NM_005034 NP_005025 5440 POLR2K 4C NM_0030938 NP_112200 81671 VMP1 4C NM_004553 NP_004544 4726 NDUFS6 4C NM_004553 NP_0055787 22980 KIAA1049 4C NM_006601 NP_006592 10728 TEBP 4C NM_007000 NP_006691 301 ANXA1	H factor 1 (complement)	HFI	3075	NP_000177	NM_000186	†C	seob0200 4C
NM_001424 NP_001415 2013 EMP2 NM_003758 NP_003749 8669 EIF3SI NM_024537 NP_078813 79587 FLJ12118 NM_153477 NP_078813 79587 FLJ12118 NM_004894 NP_004885 9556 C14or£2 NM_018322 NP_060792 55776 C6orf64 NM_002106 NP_005025 5440 POLR2K NM_005034 NP_066968 9988; 402563; 401385 DMTFI NM_030938 NP_112200 81671 VMPI NM_004553 NP_004544 4726 NDUFS6 NM_004553 NP_0055787 22980 KIAA1049 NM_006601 NP_006592 10728 TEBP	annexin Al	ANXAI	301	NP_000691	NM_000700	†C	seob0182
NIM_001424 NP_001415 2013 EMP2 NIM_003758 NP_003749 8669 EIF3S1 NM_024537 NP_078813 79587 FLJ12118 NM_024537 NP_070582 8409 UXT NM_004894 NP_060792 5576 C14orf2 NM_018322 NP_060792 55776 C6orf64 NM_002106 NP_005025 5440 POLR2K NM_005034 NP_066968 9988; DMTF1 NM_021145 NP_066968 9988; DMTF1 NM_030938 NP_112200 81671 VMP1 NM_04553 NP_004544 4726 NDUFS6 NM_014972 NP_055787 22980 KIAA1049 NM_000978 NP_00969 9349 RPL23	unactive progesterone receptor, 23 kD	TEBP	10728	NP_006592	NM_006601	†C	seob0168 4C
NM_001424 NP_001415 2013 EMP2 NM_003758 NP_003749 8669 EIF3SI NM_024537 NP_078813 79587 FLJ12118 NM_004894 NP_004885 9556 C14orf2 NM_0018322 NP_060792 55776 C6orf64 NM_002106 NP_005025 5440 POLR2K NM_005034 NP_066968 9988; 402563; 401385 DMTF1 NM_030938 NP_112200 81671 VMP1 NM_004553 NP_004544 4726 NDUFS6 NM_014972 NP_055787 22980 KIAA1049	ribosomal protein L23	RPL23	9349	NP_000969	NM_000978	#C	seob0154 4C
NM_001424 NP_001415 2013 EMP2 NM_003758 NP_003749 8669 EIF3S1 NM_003758 NP_078813 79587 FLJ12118 NM_0024537 NP_0705582 8409 UXT NM_004894 NP_060792 55776 C6orf64 NM_018322 NP_060792 55776 C6orf64 NM_002106 NP_005025 5440 POLR2K NM_005034 NP_066968 9988; DMTF1 NM_030938 NP_112200 81671 VMP1 NM_004553 NP_004544 4726 NDUFS6	KIAA1049 protein	KIAA1049	22980	NP_055787	NM_014972	†C	seob0122 4C
NM_001424 NP_001415 2013 EMP2 NM_003758 NP_003749 8669 EIF3S1 NM_003758 NP_003749 8669 EIF3S1 NM_0024537 NP_078813 79587 FLJ12118 NM_004894 NP_705582 8409 UXT NM_018322 NP_060792 5576 C14orf2 NM_002106 NP_002097 3015 H2AFZ NM_005034 NP_066968 9988; DMTF1 NM_030938 NP_112200 81671 VMP1 NM_002192 NP_002183 3624 INHBA	NADH dehydrogenase (ubiquinone) Fe-S protein 6, 13	NDUFS6	4726	NP_004544	NM_004553	4C	seob0089 4C
NM_001424 NP_001415 2013 EMP2 NM_003758 NP_003749 8669 EIF3S1 NM_024537 NP_078813 79587 FLJ12118 NM_024537 NP_705582 8409 UXT NM_004894 NP_060792 5576 C14orf2 NM_018322 NP_060792 55776 C6orf64 NM_005034 NP_005025 5440 POLR2K NM_021145 NP_066968 9988; 402563; 401385 DMTF1 NM_030938 NP_112200 81671 VMP1	inhibin, beta A (activin A, activin AB alpha polyp	INHBA	3624	NP_002183	NM_002192	4C	seob0085 4C
4C NM_001424 NP_001415 2013 EMP2 4C NM_003758 NP_003749 8669 EIF3S1 4C NM_024537 NP_078813 79587 FLJ12118 4C NM_004894 NP_004885 9556 C14ort2 4C NM_018322 NP_060792 55776 C6orf64 4C NM_002106 NP_002097 3015 H2AFZ 4C NM_005034 NP_066968 9988; DMTF1 4C NM_021145 NP_066968 9988; DMTF1	likely ortholog of rat vacuole membrane protein 1	VMPI	81671	NP_112200	NM_030938	4C	seob0065
NM_001424 NP_001415 2013 EMP2 NM_003758 NP_003749 8669 EIF3S1 NM_024537 NP_078813 79587 FLJ12118 NM_153477 NP_705582 8409 UXT NM_004894 NP_004885 9556 C14orf2 NM_018322 NP_060792 55776 C6orf64 NM_005034 NP_005025 5440 POLR2K	cyclin D binding myb-like transcription factor 1	DMTFI	9988; 402563; 401385	NP_066968	NM_021145	4C	seob0046 4C
4C NM_001424 NP_001415 2013 EMP2 4C NM_003758 NP_003749 8669 EIF3S1 4C NM_024537 NP_078813 79587 FLJ12118 4C NM_153477 NP_705582 8409 UXT 4C NM_004894 NP_004885 9556 C14orf2 4C NM_018322 NP_060792 55776 C6orf64 4C NM_002106 NP_002097 3015 H2AFZ	polymerase (RNA) II (DNA directed) polypeptide K,	POLR2K	5440	NP_005025	NM_005034	4C	seob0031
NM_001424 NP_001415 2013 EMP2 NM_003758 NP_003749 8669 EIF3S1 NM_024537 NP_078813 79587 FLJ12118 NM_153477 NP_705582 8409 UXT NM_004894 NP_004885 9556 C14orf2 NM_018322 NP_060792 55776 C6orf64	H2A histone family, member Z	H2AFZ	3015	NP_002097	NM_002106	40	seoa9935
4C NM_001424 NP_001415 2013 EMP2 4C NM_003758 NP_003749 8669 EIF3S1 4C NM_024537 NP_078813 79587 FLJ12118 4C NM_153477 NP_705582 8409 UXT 4C NM_004894 NP_004885 9556 C14orf2	chromosome 6 open reading frame 64	C6orf64	55776	NP_060792	NM_018322	4C	seoa9930 4C
4C NM_001424 NP_001415 2013 EMP2 4C NM_003758 NP_003749 8669 EIF3S1 4C NM_024537 NP_078813 79587 FLJ12118 4C NM_153477 NP_705582 8409 UXT		C14orf2	9556	NP_004885	NM_004894	4C	seoa9889
4C NM_001424 NP_001415 2013 EMP2 4C NM_003758 NP_003749 8669 EIF3S1 4C NM_024537 NP_078813 79587 FLJ12118	ubiquitously-expressed transcript	TXU	8409	NP_705582	NM_153477	4C	seoa9873
NM_001424 NP_001415 2013 EMP2 NM_003758 NP_003749 8669 EIF3S1	hypothetical protein FLJ12118	FLJ12118	79587	NP_078813	NM_024537	4C	seoa9870 4C
4C NM_001424 NP_001415 2013 EMP2 epithelial me	eukaryotic translation initiation factor 3, subuni	EIF3S1	8669	NP_003749	NM_003758	4C	seoa9855
	epithelial membrane protein 2	EMP2	2013	NP_001415	NM_001424	4C	seoa9828
NM_080832 NP_543022 140886 PABPC5 poly(A) bind	poly(A) binding protein, cytoplasmic 5	PABPC5	140886	NP_543022	NM_080832	4C	seoa9817 4C

heat shock 70kDa protein 5 (glucose-regulated prot	HSPA5 h	3309	NP_005338	INIM_UUDDA4/		30001131
DnaJ (Hsp40) homolog, subfamily C, member 8	DNAJC8 [6	NP_055095	NM 005347	4C +C	
ATP synthase, H+ transporting, mitochondrial F0 co	ATP5I A		NP_009031	00/100 MN	A) &	sco01143
transmembrane protein 14B	TMEM14B to	81853	NP_112231	NM_030969		
N-myc (and STAT) interactor	NMI	9111	NP_004679	NM_004688		seop1093 4C
aryl hydrocarbon receptor nuclear translocator-lik	ARNTL2 a	56938	NP_064568	NM_020183		Seob1037 4C
autophagy Apg3p/Aut1p-like	APG3 2	64422	NP_071933	NM_022488		SE001032 4C
stress-associated endoplasmic reticulum protein 1	SERP1 s	27230	NP_055260	NM_014445		Sc001000 4C
polymerase (RNA) II (DNA directed) polypeptide G	POLR2G I	5436	NP_002687	NM_002696		Se0000/2 4C
pyruvate dehydrogenase kinase, isoenzyme 1	PDK1	5163	NP_002601	NM_002610	45	Se000840
mitochondrial ribosomal protein S25	MRPS25	64432	NP_071942	NM_022497	÷ (36000031
G. Somoto, I. (Barcotti		57026; 23616				0000001
lectin, galactoside-binding, soluble, 1 (galectin	LGALSI	3956;	NP_002296	NM_002305	4C	seob0810 4C
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide	RIG-I	23586	NP_055129	NM_014314	40	seobo /os
survival of motor neuron protein interacting prote	SIP1	8487	NP_003607	NM_003616	6	
proline-rich nuclear receptor coactivator	PNRCI	10957	NP_006804	NM_006813	46	2000/43/40
CDC16 cell division cycle 16 homolog (S. cerevisia	CDC16	8881	NP_003894	NM_003903	6	seop0/03/4C
apoptosis related protein APR-3	3-Apr	51374	NP_542159	NM_016085	4C	seob0514 4C
HSPCO34 protein	LOC51668	51668	NP_057210	NM_016126	4C	seob0497/4C
karyopherin alpha 4 (importin alpha 3)	KPNA4	3840	NP_002259	NM_002268	4C	Seob04 /8 4C
ubiquitous tetratricopeptide containing protein Ro	RoXaN	23264	NP_060060	NM_017590	40	seob0466 4C
ubiquitin specific protease 38	USP38	84640	NP_115946	NM_032557	÷	Se000418
wingless-type MMTV integration site family, member	WNT5A	7474	NP_003383	NM_003392	4C	seob0376 4C
nucleolar protein 7, 27kDa	NOL7	51406	NP_057251	NM_016167	4C	seob0321
chemokine (C-C motif) ligand 13	CCL13	6357	NP_005399	NM_005408	4C	seob0308 4C

ADP-ribosylation factor-like 10C	ARL10C	55207	NP_060654	NM_018184	seob1906 4C	seob19
histamine receptor H1	HRHI	3269	NP_000852	NM_000861	seob1879 4C	seob18
Williams-Beuren syndrome chromosome region 1	WBSCRI	7458	NP_114381	NM_022170	seob1862 4C	seobl
geminin, DNA replication inhibitor	GMNN	51053	NP_056979	NM_015895	seob1834 4C	seobl
dynein, cytoplasmic, light intermediate polypeptid	DNCLI2	1783	NP_006132	NM_006141	seob1793 4C	seobl
cytoplasmic FMR1 interacting protein 1	CYFIPI	23191	NP_055423	NM_014608	seob1770 4C	seobl
				NM_018572	seob1766 4C	seob1
mitochondrial ribosomal protein L37	MRPL37	51253	NP_057575	NM_016491	seob1748 4C	seob1
molecule possessing ankyrin repeats induced by lip	MAIL	64332	NP_113607	NM_031419	seob1746 4C	seob1
EBNA1 binding protein 2	EBNA1BP2	10969	NP_006815	NM_006824	seob1737 4C	seob1
mitochondrial ribosomal protein L20	MRPL20	55052	NP_060441	NM_017971	seob1667 4C	seobl
chromosome 4 open reading frame 15	C4orf15	79441	NP_078787	NM_024511	seob1660 4C	seobl
zinc finger protein 533	ZNF533	151126	NP_689733	NM_152520	seob1646 4C	seobl
LanC lantibiotic synthetase component C-like 1 (ba	LANCLI	10314	NP_006046	NM_006055	seob1617 4C	seobl
synovial sarcoma translocation gene on chromosome	SS18L2	51188	NP_057389	NM_016305	seob1574 4C	seobl
trophoblast glycoprotein	TPBG	7162	NP_006661	NM_006670	seob1426 4C	seobl
MUF1 protein	MUFI	10489	NP_006360	NM_006369	seob1420 4C	seob1
eukaryotic translation elongation factor 1 epsilon	EEF1E1	9521	NP_004271	NM_004280	1414 4C	seob1414
				NM_014949	1385 4C	seob1385
transcription factor CP2	TFCP2	7024	NP_005644	NM_005653	1383 4C	seob1383
v-fos FBJ murine osteosarcoma viral oncogene homol	FOS	2353	NP_005243	NM_005252	1322 4C	seob1322
reticulon 4	RTN4	57142	NP_997404	NM_020532	seob1319 4C	seobl
hypothetical protein FLJ14753	FLJ14753	84641	NP_115947	NM_032558	seob1318 4C	seob l
solute carrier family 1 (glutamate/neutral amino a	SLC1A4	6509	NP_003029	NM_003038	seob1243 4C	seobl
ARP3 actin-related protein 3 homolog (yeast)	ACTR3	10096	NP_005712	NM_005721	seob1231 4C	seobl

NP_057581 51258 NP_002262 3843 NP_004371 1387 NP_002088 2971 NP_000282 5230 NP_689582 126969 NP_003324 7311 NP_058523 5937 NP_060922 55836 NP_003378 7456	NM_001315 NM_001315 NM_001315 NM_017824 NM_016497 NM_002271 NM_002297 NM_002097 NM_000291 NM_0152369 NM_0152369 NM_018452 NM_018452 NM_018452 NM_003387 NM_003387	seob2185 4C seob2188 4C seob2221 4C seob2642 4C seob2658 4C seob2750 4C seob2797 4C seob2803 4C seob2807 4C seob2810 4C
NP_057581 51258 MRPL51 NP_002262 3843 KPNB3 NP_004371 1387 CREBBP NP_002088 2971 GTF3A NP_000282 5230 PGK1 NP_689582 126969 MGC45474 NP_003324 7311 UBA52 NP_058523 5937 RBMS1 NP_060922 55836 C6orf35		seob2185 4C seob2188 4C seob2221 4C seob2642 4C seob2658 4C seob2697 4C seob2750 4C seob2797 4C seob2803 4C seob2801 4C
NP_057581 51258 MRPL51 NP_002262 3843 KPNB3 NP_004371 1387 CREBBP NP_002088 2971 GTF3A NP_000282 5230 PGK1 NP_689582 126969 MGC45474 NP_003324 7311 UBA52 NP_058523 5937 RBMS1		seob2185 4C seob2188 4C seob2221 4C seob2642 4C seob2658 4C seob2697 4C seob2790 4C seob2791 4C seob2803 4C
NP_057581 51258 MRPL51 NP_002262 3843 KPNB3 NP_004371 1387 CREBBP NP_002088 2971 GTF3A NP_000282 5230 PGK1 NP_689582 126969 MGC45474 NP_003324 7311 UBA52		seob2185 4C seob2188 4C seob2221 4C seob2642 4C seob2658 4C seob2697 4C seob2750 4C seob2797 4C
NP_057581 51258 MRPL51 NP_002262 3843 KPNB3 NP_004371 1387 CREBBP NP_002088 2971 GTF3A NP_000282 5230 PGK1 NP_689582 126969 MGC45474		seob2185 4C seob2188 4C seob2221 4C seob2642 4C seob2658 4C seob2697 4C seob2750 4C
NP_057581 51258 MRPL51 NP_002262 3843 KPNB3 NP_004371 1387 CREBBP NP_002088 2971 GTF3A NP_000282 5230 PGK1		seob2185 4C seob2188 4C seob2221 4C seob2642 4C seob2658 4C seob2697 4C seob2750 4C
NP_057581 51258 MRPL51 NP_002262 3843 KPNB3 NP_004371 1387 CREBBP NP_002088 2971 GTF3A		seob2185 4C seob2188 4C seob2221 4C seob2642 4C seob2658 4C seob2697 4C
NP_057581 51258 MRPL51 NP_002262 3843 KPNB3 NP_004371 1387 CREBBP		seob2185 4C seob2188 4C seob2221 4C seob2642 4C seob2658 4C
NP_057581 51258 MRPL51 mitochondrial NP_002262 3843 KPNB3 karyopherin (i		seob2185 4C seob2188 4C seob2221 4C seob2642 4C
NP_057581 51258 MRPL51 mitochondrial		seob2185 4C seob2188 4C seob2221 4C
		seob2185 4C seob2188 4C
NP 060294 54708 FLJ20445		seob2185 4C
5 NP_620583 1432 MAPK14 mitogen-activated protein kinase 14		
NP_758961 1997 ELF1 E74-like factor 1 (ets domain transcription factor		seob2169 4C
NP_054872 29066 ZC3HDC7 zinc finger CCCH type domain containing 7	NM_014153	seob2148 4C
NP_000167 2908 NR3C1 nuclear receptor subfamily 3, group C, member 1 (g	NM_000176	seob2139 4C
NP_067006 57826 RAP2C RAP2C, member of RAS oncogene family	NM_021183	seob2134 4C
NP_057671 11335 CBX3 chromobox homolog 3 (HP1 gamma homolog, Drosophila	NM_016587	seob2108 4C
NP_009209 11337; GABARAP GABA(A) receptor-associated protein 23399;	NM_007278	seob2081 4C
NP_002786 5691 PSMB3 proteasome (prosome, macropain) subunit, beta type	NM_002795	seob2077 4C
NP_003190 6925 TCF4 transcription factor 4	NM_003199	seob2011 4C
NP_842565 6711; 56969 SPTBN1 spectrin, beta, non-erythrocytic 1	NM_003128	seob1972 4C
NP_006760 8543 LMO4 LIM domain only 4	NM_006769	seob1916 4C
NP_115714 84298 MGC14817 hypothetical protein MGC14817	NM_032338	seob1911 4C
NP_009089 7812 DISI55E NRAS-related gene	NM_007158	seob1908 4C

				XM_088391	4C	seob3462 4C
t-complex-associated-testis-expressed 1-like 1	TCTELI	6993	NP_006510	NM_006519	4C	seob3455
selenoprotein T	SELT	51714; 389704	NP_057359	NM_016275	4C	seob3451
MAM domain containing 2	MAMDC2	256691	NP_694999	NM_153267	4C	seob3367_4C
RAP1B, member of RAS oncogene family	RAPIB	5908	NP_056461	NM_015646	4C	seob3356_4C
ribosomal protein, large P2	RPLP2	1819	NP_000995	NM_001004	4C	seob3326 4C
ubiquitin-conjugating enzyme E2 variant 1	UBE2V1	7335; 387521; 387522	NP_954595	NM_199203	4C	seob3313
transforming growth factor, beta-induced, 68kDa	TGFBI	7045	NP_000349	NM_000358	4C	seob3307
stromal antigen 2	STAG2	10735	NP_006594	NM_006603	4C	seob3303
microtubule-actin crosslinking factor 1	MACFI	23499	NP_149033	NM_033044	4C	seob3191
chromosome 14 open reading frame 2	C14orf2	9556	NP_004885	NM_004894	4C	seob3189
KIAA0372	KIAA0372	9652	NP_055454	NM_014639	4C	seob3182
ectonucleotide pyrophosphatase/phosphodiesterase 1	ENPP1	5167	NP_006199	NM_006208	4C	seob3170 4C
chromosome X open reading frame 6	CXorf6	10046	NP_005482	NM_005491	4C	seob3169 4C
chromosome 10 open reading frame 86	C10orf86	54780	NP_060085	NM_017615	4C	seob3154_4C
HSPC056 protein	HSPC056	25852	NP_998819	NM_015396	4C	seob3119
homeodomain interacting protein kinase 3	HIPK3	10114	NP_005725	NM_005734	4C	seob3076
CDK2-associated protein 1	CDK2AP1	8099	NP_004633	NM_004642	4C	seob3064
hypothetical protein FLJ10970	FLJ10970	55273	NP_060756	NM_018286	4C	seob3007
praja 2, RING-H2 motif containing	PJA2	9867	NP_055634	NM_014819	4C	seob2994 4C
CAAX box 1	CXXI	8933	NP_003919	NM_003928	4C	seob2987 4C
protein phosphatase 1, regulatory (inhibitor) subu	PPP1R12A	4659	NP_002471	NM_002480	4C	seob2966 4C
ribosomal protein S19	RPS19	6223	NP_001013	NM_001022	4C	seob2959 4C
interleukin enhancer binding factor 2, 45kDa	ILF2	3608	NP_004506	NM_004515	4C	seob2936 4C

H3 histone, family 3A	H3F3A	3020	NP_002098	NM_002107	4C	seob4122
ubiquinol-cytochrome c reductase complex (7.2 kD)	HSPC051	29796	NP_037519	NM_013387	†	seob4118 4C
potassium channel modulatory factor 1	KCMF1	56888	NP_064507	NM_020122	†C	seob4108 4C
RAD23 homolog B (S. cerevisiae)	RAD23B	5887	NP_002865	NM_002874	‡C	seob4075 4C
lactate dehydrogenase A	LDHA	3939	NP_005557	NM_005566	4C	seob4050 4C
myeloid cell leukemia sequence 1 (BCL2-related)	MCL1	4170	NP_877495	NM_021960	4C	seob4039 4C
				NM_153012	4C	seob4030 4C
CP110 protein	CP110	9738	NP_055526	NM_014711	1C	seob4029 4C
CGI-72 protein	CGI-72	51105	NP_940915	NM_016018	4C	seob3923 4C
oxysterol binding protein-like 8	OSBPL8	114882	NP_065892	NM_020841	4C	seob3887
sarcoglycan, epsilon	SGCE	8910	NP_003910	NM_003919	4C	seob3869
guanine nucleotide binding protein (G protein), al	GNAI3	2773	NP_006487	NM_006496	4C	seob3854
chromosome 14 open reading frame 11	C14orf11	55837	NP_060923	NM_018453	4C	seob3751
small nuclear ribonucleoprotein D2 polypeptide 16.	SNRPD2	6633	NP_808210	NM_004597	†)	seob3734 4C
nucleophosmin (nucleolar phosphoprotein B23, numat	NPM1	4869	NP_954654	NM_002520	‡C	seob3731 4C
SON DNA binding protein	SON	6651	NP_620305	NM_058183	ţ.	seob3699 4C
CD14 antigen	CD14	929; 4695	NP_000582	NM_002488	ţ.	seob3670 4C
NCK-associated protein 1	NCKAPI	10787	NP_995314	NM_013436	t)	seob3533 4C
RAB4A, member RAS oncogene family	RAB4A	5867; 10638 RAB4A	NP_004569	NM_004578	Ċ	seob3520 4C
numb homolog (Drosophila)	NUMB	8650	NP_003735	NM_003744	4C	seob3518
ribosomal protein, large, P1	RPLP1	6176	NP_998890	NM_001003	4C	seob3513
actin related protein 2/3 complex, subunit 2, 34kD	ARPC2	10109	NP_690601	NM_152862	4C	seob3499
down-regulator of transcription 1, TBP-binding (ne	DR1	1810	NP_001929	NM_001938	4C	seob3485
CDA11 protein	CDA11	83940	NP_114415	NM_032026	4C	seob3468
dynein, cytoplasmic, intermediate polypeptide 2	DNC12	1781	NP_001369	NM_001378	ţ,	seob3464 4C

ubiquitin specific protease 8	USP8 ul	9101	NP_005145	NM_005154	4C	seob4835 4C
splicing factor, arginine/serine-rich 10 (transfor	SFRS10 sp	6434	NP_004584		4C	seob4807
solute carrier family 38, member 2	SLC38A2 sc	54407	NP_061849	NM_018976	4C	seob4766
mortality factor 4 like 1	MORF4L1 m	10933	NP_996670	NM_206839	4C	seob4752
N-myc downstream regulated gene 1	NDRG1 N	10397	NP_006087	NM_006096	4C	seob4726
karyopherin (importin) beta 1	KPNB1 k	3837	NP_002256	NM_002265	4C	seob4669 4C
tryptophan rich basic protein	WRB tr	7485	NP_004618	NM_004627	4C	seob4621 4C
protein phosphatase 1, regulatory (inhibitor) subu	PPP1R2 p	5504	NP_006232	NM_006241	4C	seob4612 4C
NADH dehydrogenase (ubiquinone) 1 beta subcomplex,	NDUFB2 N	4708	NP_004537	NM_004546	4C	seob4579 4C
mannose-6-phosphate receptor binding protein 1	M6PRBP1 m	10226	NP_005808	NM_005817	4C	seob4570 4C
fibroblast growth factor receptor 2 (bacteria-expr	FGFR2 fi	2263	NP_075420	NM_023028	4C	seob4555
gamma-SNAP-associated factor 1	GAF1 g	26056	NP_056285	NM_015470	4C	seob4545
eukaryotic translation initiation factor 2B, subun	EIF2B3 e	8891	NP_065098	NM_020365	4C	seob4539 4C
ribosomal protein S27-like	RPS27L ri	51065	NP_057004	NM_015920	4C	seob4515
glycogenin	GYG g	2992	NP_004121	NM_004130	4C	seob4492 4C
Fc fragment of lgG, low affinity IIa, receptor for	FCGR2A F	2212	NP_067674	NM_021642	4C	seob4480 4C
oxysterol binding protein-like 2	OSBPL2 0	9885	NP_653081	NM_144498	4C	seob4427 4C
sin3-associated polypeptide, 18kDa	SAP18 si	10284	NP_005861	NM_005870	4C	seob4419 4C
leucine zipper transcription factor-like 1	LZTFL1	54585	NP_065080	NM_020347	4C	seob4333
ribosomal protein S25	RPS25 ri	6230	NP_001019	NM_001028	4C	seob4303
RNA polymerase II associated protein 1	RPAP1 R	26015	NP_056355	NM_015540	4C	seob4273
phospholipase A2, group IVA (cytosolic, calcium-de	PLA2G4A p	5321	NP_077734	NM_024420	4C	seob4213
prolylcarboxypeptidase (angiotensinase C)	PRCP p	5547; 360219	NP_955450	NM_199418	4C	seob4209 4C
hypothetical protein LOC283820	LOC283820 h	283820; 23420	NP_775885	NM_014287	4C	seob4197 4C
LSM6 homolog, U6 small nuclear RNA associated (S.	LSM6 L	11157	110600 ⁻ dN	NM_007080	4C	seob4191 4C

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NM_001386 NP_001377 1808 DPYSL2 3 NM_025134 NP_001377 1808 DPYSL2 3 NM_025134 NP_079410 80205; FLJ12178 4 NM_025134 NP_079410 80205; FLJ12178 5 NM_01402 NP_001393 1915; EEF1A1 3 NM_01402 NP_00373 389411 29894 0 NM_003273 NP_115649 84233 DKFZp586C1 0 NM_016613 NP_0037470 8705; 6222 B3GALT4 0 NM_013338 NP_057697 51313 DKFZp434L1 0 NM_144684 NP_0573285 147657 MGC32104 0 NM_015577 NP_062365 56288 PARD3 0 NM_005470 NP_056392 26064 RAH14 0 NM_005496 NP_055330 2388 KLHDC2 0 NM_001341 NP_005487 10051 SMC4L1 0 NM_0013416 NP_005335 1603<	RAB8B, member RAS oncogene family	RAB8B	51762	NP_057614	NM_016530	4	seono 660
NM_001386 NP_001377 1808 DPYSL2 3 NM_025134 NP_001377 1808 DPYSL2 3 NM_001402 NP_001393 1915; EEF1A1 4 NM_006965 NP_008896 7572 ZNF24 5 NM_006965 NP_008896 7572 ZNF24 6 NM_006965 NP_008896 7572 ZNF24 7 NM_0032273 NP_115649 84233 DKFZp586C1 8 NM_016613 NP_003773 8705; 6222 B3GALT4 8 NM_016613 NP_037470 29880 ALG5 8 NM_0144684 NP_037470 29880 ALG5 8 NM_015577 NP_053285 147657 MGC32104 8 NM_015577 NP_056392 26064 RAH14 8 NM_005470 NP_056392 26064 RAH14 8 NM_005470 NP_056392 26064 RAH14 8 NM_005470 NP_055130 23588 KLHDC2 <	TRIAD3 protein	TRIAD3		NP_996999	NM_207111	4C	
NM_001386 NP_001377 1808 DPYSL2 NM_025134 NP_079410 80205; FLJ12178 NM_025134 NP_079410 80205; FLJ12178 NM_025134 NP_079410 80205; FLJ12178 NM_02666 NP_001393 1915; EEF1A1 NM_006965 NP_008896 7572 ZNF24 NM_006965 NP_003773 8705; 6222 B3GALT4 NM_0032273 NP_003773 8705; 6222 B3GALT4 NM_016613 NP_037470 29880 ALG5 NM_0144684 NP_053285 147657 MGC32104 NM_015577 NP_056392 26064 RAI14 NM_005470 NP_056392 26064 RAI14 NM_00264 NP_0056392 26064 RAI14 NM_005496 NP_005487 10051 SMC4L1 NM_001344 NP_005330 23588 KLHDC2 NM_0012168 NP_036300 26332 FBXO2					NM_031268	tc C	seob5748 4C
NM_001386 NP_001377 1808 DPYSL2 NM_0025134 NP_001377 1808 DPYSL2 NM_0025134 NP_001393 1915; EEF1A1 NM_001402 NP_001393 1915; EEF1A1 NM_006965 NP_001393 1915; EEF1A1 NM_0032273 NP_115649 84233 DKFZp586C1 NM_032273 NP_003773 8705; 6222 B3GALT4 NM_016613 NP_057697 51313 DKFZp586C1 NM_013338 NP_037470 29880 ALG5 NM_015577 NP_062565 56288 PARD3 NM_0173552 NP_775823 205428 MGC33165 NM_005496 NP_005487 10051 SMC4L1 NM_001344 NP_0055130 23588 KLHDC2 NM_005749 NP_005740 10140 TOB1	F-box only protein 2	FBXO2		NP_036300	NM_012168	tC	seob5743 4C
NM_001386 NP_001377 1808 DPYSL2 NM_025134 NP_001377 1808 DPYSL2 NM_025134 NP_001393 1915; EEF1A1 NM_001402 NP_001393 1915; EEF1A1 NM_006965 NP_001393 1915; EEF1A1 NM_006965 NP_008896 7572 ZNF24 NM_0032273 NP_115649 84233 DKFZp586C1 NM_016613 NP_003773 8705; 6222 B3GALT4 NM_013338 NP_057697 51313 DKFZp434L1 NM_013338 NP_037470 29880 ALG5 NM_019619 NP_053285 147657 MGC32104 NM_005470 NP_056392 26064 RAH14 NM_005496 NP_775823 205428 MGC33365 NM_005496 NP_005487 10051 SMC4L1 NM_01341 NP_0055130 23588 KLHDC2 NM_001344 NP_001335 1603 DAD1	transducer of ERBB2, 1	TOBI		NP_005740	NM_005749	†C	seob5726 4C
NM_001386 NP_001377 1808 DPYSL2 NM_025134 NP_079410 80205; FLJ12178 NM_001402 NP_001393 1915; EEF1AI NM_006965 NP_008896 7572 ZNF24 NM_0032273 NP_115649 84233 DKFZp586C1 NM_016613 NP_057697 51313 DKFZp586C1 NM_013338 NP_057697 51313 DKFZp434L1 NM_019619 NP_0537470 29880 ALG5 NM_019619 NP_062565 56288 PARD3 NM_005470 NP_053285 147657 MGC32104 NM_005470 NP_056392 26064 RAH14 NM_005496 NP_005461 10006 ABH NM_005496 NP_005487 10051 SMC4L1 NM_014315 NP_055130 23588 KLHDC2	defender against cell death 1	DADI		NP_001335	NM_001344	4C	seob5645
NM_001386 NP_001377 1808 DPYSL2 NM_0025134 NP_001393 1808 DPYSL2 NM_001402 NP_001393 1915; EEF1A1 NM_006965 NP_008896 7572 ZNF24 NM_0032273 NP_115649 84233 DKFZp586C1 NM_016613 NP_003773 8705; 6222 B3GALT4 NM_013338 NP_057697 51313 DKFZp434L1 NM_0144684 NP_053285 147657 MGC32104 NM_015577 NP_062565 56288 PARD3 NM_005470 NP_056392 26064 RA114 NM_005496 NP_775823 205428 MGC33365 NM_005496 NP_005487 10051 SMC4L1	kelch domain containing 2	KLHDC2		NP_055130	NM_014315	4C	seob5562
NM_001386 NP_001377 1808 DPYSL2 NM_025134 NP_079410 80205; FLJ12178 NM_001402 NP_001393 1915; EEF1A1 NM_006965 NP_001393 1915; EEF1A1 NM_006965 NP_008896 7572 ZNF24 NM_0032273 NP_115649 84233 DKFZp586C1 NM_016613 NP_003773 8705; 6222 B3GALT4 NM_013338 NP_057697 51313 DKFZp434L1 NM_013338 NP_037470 29880 ALG5 NM_015577 NP_0653285 147657 MGC32104 NM_005470 NP_056392 26064 RAH14 NM_002064 NP_775823 205428 MGC33365 NM_002064 NP_002055 2745 GLRX	SMC4 structural maintenance of chromosomes 4-like	SMC4L1		NP_005487	NM_005496	4C	seob5523
NM_001386 NP_001377 1808 DPYSL2 NM_025134 NP_079410 80205; FLJ12178 NM_001402 NP_001393 1915; EEF1A1 NM_006965 NP_008896 7572 ZNF24 NM_032273 NP_115649 84233 DKFZp586C1 NM_022551 NP_003773 8705; 6222 B3GALT4 NM_013338 NP_057697 51313 DKFZp434L1 NM_144684 NP_053285 147657 MGC32104 NM_015577 NP_062565 56288 PARD3 NM_005470 NP_0056392 26064 RAH14 NM_173552 NP_775823 205428 MGC33365	glutaredoxin (thioltransferase)	GLRX		NP_002055	NM_002064	4C	seob5490
NM_001386 NP_001377 1808 DPYSL2 NM_001402 NP_079410 80205; FLJ12178 NM_001402 NP_001393 1915; EEF1A1 NM_006965 NP_008896 7572 ZNF24 NM_032273 NP_115649 84233 DKFZp586C1 NM_016613 NP_003773 8705; 6222 B3GALT4 NM_013338 NP_037470 29880 ALG5 NM_019619 NP_053285 147657 MGC32104 NM_015577 NP_056392 26064 RAI14 NM_005470 NP_005461 10006 ABI1	hypothetical protein MGC33365	MGC33365		NP_775823	NM_173552	4C	seob5458
NM_001386 NP_001377 1808 DPYSL2 NM_025134 NP_079410 80205; FLJ12178 NM_001402 NP_001393 1915; EEF1A1 NM_006965 NP_008896 7572 ZNF24 NM_032273 NP_115649 84233 DKFZp586C1 NM_022551 NP_003773 8705; 6222 B3GALT4 NM_013338 NP_057697 51313 DKFZp434L1 NM_144684 NP_653285 147657 MGC32104 NM_015577 NP_056392 26064 RAH4	abl-interactor l	ABH		NP_005461	NM_005470	4C	seob5441
NM_001386 NP_001377 1808 DPYSL2 NM_025134 NP_079410 80205; FLJ12178 NM_001402 NP_001393 1915; EEF1A1 NM_006965 NP_008896 7572 ZNF24 NM_032273 NP_115649 84233 DKFZp586C1 NM_016613 NP_003773 8705; 6222 B3GALT4 NM_013338 NP_037470 29880 ALG5 NM_019619 NP_062565 56288 PARD3		RAH4	26064	NP_056392	NM_015577	4C	seob5319
NM_001386 NP_001377 1808 DPYSL2 NM_025134 NP_079410 80205; FLJ12178 NM_001402 NP_001393 1915; EEF1A1 NM_006965 NP_008896 7572 ZNF24 NM_032273 NP_115649 84233 DKFZp586C1 NM_022551 NP_003773 8705; 6222 B3GALT4 NM_016613 NP_057697 51313 DKFZp434L1 NM_013338 NP_037470 29880 ALG5 NM_144684 NP_653285 147657 MGC32104	par-3 partitioning defective 3 homolog (C. elegans	PARD3	56288	NP_062565	NM_019619	4C	seob5240 4C
NM_001386 NP_001377 1808 DPYSL2 NM_025134 NP_079410 80205; FLJ12178 NM_001402 NP_001393 1915; EEF1A1 NM_006965 NP_008896 7572 ZNF24 NM_032273 NP_115649 84233 DKFZp586C1 NM_022551 NP_003773 8705; 6222 B3GALT4 NM_016613 NP_057697 51313 DKFZp434L1 NM_013338 NP_037470 29880 ALG5	hypothetical protein MGC32104	MGC32104	147657	NP_653285	NM_144684	4C	seob5219 4C
NM_001386 NP_001377 1808 DPYSL2 NM_025134 NP_079410 80205; FLJ12178 NM_001402 NP_001393 1915; EEF1A1 NM_006965 NP_008896 7572 ZNF24 NM_032273 NP_115649 84233 DKFZp586C1 NM_022551 NP_003773 8705; 6222 B3GALT4 NM_016613 NP_057697 51313 DKFZp434L1	asparagine-linked glycosylation 5 homolog (yeast,	ALG5		NP_037470	NM_013338	4C	seob5213
NM_001386 NP_001377 1808 DPYSL2 NM_025134 NP_079410 80205; FLJ12178 29962 NM_001402 NP_001393 1915; EEF1A1 NM_006965 NP_008896 7572 ZNF24 NM_032273 NP_115649 84233 DKFZp586C1 NM_022551 NP_003773 8705; 6222 B3GALT4		DKFZp434L1 42	51313	NP_057697	NM_016613	4C	seob5174 4C
NM_001386 NP_001377 1808 DPYSL2 NM_025134 NP_079410 80205; FLJ12178 29962 29962 FLJ12178 NM_001402 NP_001393 1915; EEF1A1 NM_006965 NP_008896 7572 ZNF24 NM_032273 NP_115649 84233 DKFZp586C1	UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase,	B3GALT4	8705; 6222	NP_003773	NM_022551	4C	seob5044 4C
NM_001386 NP_001377 1808 DPYSL2 NM_025134 NP_079410 80205; FLJ12178 NM_001402 NP_001393 1915; EEF1A1 NM_006965 NP_008896 7572 ZNF24	hypothetical	DKFZp586C1 924	84233	NP_115649	NM_032273	4C	seob5032
NM_001386 NP_001377 1808 DPYSL2 NM_025134 NP_079410 80205; FLJ12178 NM_001402 NP_001393 1915; EEF1A1 389411 EEF1A1	zinc finger protein 24 (KOX 17)	ZNF24	7572	NP_008896	NM_006965	4C	seob5012
NM_001386	eukaryotic translation elongation factor 1 alpha 1	EEFIAI	1915; 389411	NP_001393	NM_001402	4C	seob5004 4C
NM_001386	hypothetical protein FLJ12178	FLJ12178	80205; 29962	NP_079410	NM_025134	4C	seob4992 4C
211 (20120)	dihydropyrimidinase-like 2	DPYSL2	1808	NP_001377	NM_001386	4C	seob4972 4C
4C NM_001560 NP_001551 3597 IL13RA1 interleukin 13 receptor, alpha 1	interleukin 13 receptor, alpha 1	IL13RA1	3597	NP_001551	NM_001560	4C	seob4945 4C
4C NM_003256 NP_003247 7079 TIMP4 tissue inhibitor of metalloproteinase 4	tissue inhibitor of metalloproteinase 4	TIMP4	7079	NP_003247	NM_003256	4C	seob4896 4C

seob5927 4C	4C	NM_024561	NP 078837	79612	FLJ22054	hynothetical protein El 1990s/
seob5954 4C	4C	NM_001304	NP_001295	1362	CPD	carboxypeptidase D
seob6004 4C	4C	NM_016053	NP_057137	51019	CGI-116	CGI-116 protein
seob6015	4C	NM_004902	NP_909132	9584	RNPC2	RNA-binding region (RNPI, RRM) containing 2
seob6041 4C	4C	NM_001949	NP_001940	1871	E2F3	E2F transcription factor 3
seob6096 4C	40	NM_002223	NP_002214	3709	ITPR2	inositol 1,4,5-triphosphate receptor, type 2
seob6131 4C	4C	NM_002916	NP_853551	5984	RFC4	replication factor C (activator 1) 4, 37kDa
seob6133	4C	NM_002298	NP_002289	3936	LCP1	lymphocyte cytosolic protein 1 (L-plastin)
seob6189 4C	4C	NM_002004	NP_001995	2224	FDPS	farnesyl diphosphate synthase (farnesyl pyrophosph
seob6206 4C	4C	NM_145702	NP_004837	9470; 200765	EIF4EL3	eukaryotic translation initiation factor 4E-like 3
seob6229 4C	4C	NM_020192	NP_064577	57002	C7orf36	chromosome 7 open reading frame 36
seob6272 4C	4C	NM_002767	NP_002758	5636	PRPSAP2	phosphoribosyl pyrophosphate synthetase-associated
seob6279 4C	4C	NM_006818	NP_006809	10962	AFIQ	ALL1-fused gene from chromosome 1q
seob6368 4C	4C	NM_001311	NP_001303	1397; 1396	CRIP2	cysteine-rich protein 2
seob6413 4C	4C	NM_016357	NP_057441	51474	EPLIN	epithelial protein lost in neoplasm beta
seob6446 4C	4C	XM_051200		79068	FTO	fatso
seob6467	4C	NM_006425	NP_006416	10569	SLU7	step II splicing factor SLU7
	4C	NM_016578	NP_057662	51773	HBXAP	hepatitis B virus x associated protein
seob6492	4C	NM_001540	NP_001531	3315	HSPB1	heat shock 27kDa protein 1
seob6525 4C	4C	NM_131105				
seob6558 4C	4C	NM_016096	NP_057180	51123	LOC51123	HSPC038 protein
seob6560 4C		NM_004229	NP_004220	9282	CRSP2	cofactor required for Sp1 transcriptional activati
		NM_019026	NP_061899	54499	LOC54499	putative membrane protein
seob6582	4C	NM_002124	NP_072049	3125; 3127; 1 3126; 31	HLA-DRB3	major histocompatibility complex, class II, DR bet
seobbb /0 4C		NM_000122	NP_000113	2071	ERCC3	excision repair cross-complementing rodent repair
VICE TOUGHT	•					

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ribosomal protein L17	RPL17	6139	NP_000976	NM_000985	4C	seob8311
ecotropic viral integration site 2A	EVI2A	2123	NP_055025	NM_014210	4C	seob8255
periostin, osteoblast specific factor	POSTN	10631	NP_006466	NM_006475	4C	seob8204 4C
roundabout, axon guidance receptor, homolog 1 (Dro	ROBO1	6091	NP_598334	NM_133631	4C	seob8104 4C
chromosome 11 open reading frame 10	Cllorf10	746	NP_055021	NM_014206	4C	seob8099 4C
poly(A) binding protein, cytoplasmic 1	PABPCI	26986	NP_002559	NM_002568	4C	seob8065
upregulated during skeletal muscle growth 5	USMG5	84833	NP_116136	NM_032747	4C	seob7981 4C
mortality factor 4 like 1	MORF4L1	10933	NP_996670	NM_206839	4C	seob7946
growth hormone inducible transmembrane protein	GHITM	27069	NP_055209	NM_014394	4C	seob7764 4C
SON DNA binding protein	SON	6651	NP_620305	NM_058183	4C	seob7744 4C
thioredoxin	TXN	7295	NP_003320	NM_003329	4C	seob7729 4C
proteasome (prosome, macropain) activator subunit	PSME4	23198	NP_055429	NM_014614	4C	seob7658 4C
ATP synthase, H+ transporting, mitochondrial F0 co	ATP5I	521	NP_009031	NM_007100	4C	seob7622 4C
S-phase kinase-associated protein 1A (p19A)	SKPIA	6500	NP_733779	NM_006930	4C	seob7584 4C
maternal G10 transcript	G10	8896	NP_003901	NM_003910	4C	seob7474 4C
polypyrimidine tract binding protein 1	PTBP1	5725	NP_787041	NM_002819	4C	seob7465 4C
CGI-121 protein	CGI-121	51002	NP_057142	NM_016058	4C	seob7432 4C
ADP-ribosylation factor domain protein 1, 64kDa	ARFDI	373	NP_150231	NM_001656	4C	seob7404 4C
DEAD (Asp-Glu-Ala-Asp) box polypeptide 24	DDX24	57062	NP_065147	NM_020414	4C	seob6853
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex	NDUFA5	4698	NP_004991	NM_005000	4C	seob6851
ATP-binding cassette, sub-family E (OABP), member	ABCEI	6059	NP_002931	NM_002940	4C	seob6844 4C
X-box binding protein 1	ХВР1	7494	NP_005071	NM_005080	4C	seob6836 4C
				NM_014883	4C	seob6835
Tu translation elongation factor, mitochondrial	TUFM	7284	NP_003312	NM_003321	4C	seob6751 4C
damage-specific DNA binding protein 1, 127kDa	DDB1	1642	NP_001914	NM_001923	4C	seob6680 4C

chromosome 2 open reading frame 11	C2orf11	130132	NP_653230	NM_144629	4C	seob9772 4C
ring finger protein 111	RNF111	54778	NP_060080	NM_017610	4C	seob9756 4C
ubiquitin specific protease 38	USP38	84640	NP_115946	NM_032557	4C	seob9750 4C
golgi phosphoprotein 2	GOLPH2	51280	NP_808800	NM_016548	4C	seob9649 4C
casein kinase 2, beta polypeptide	CSNK2B	1460	NP_001311	NM_001320	4C	seob9645 4C
proteasome (prosome, macropain) subunit, alpha typ	PSMA4	5685	NP_002780	NM_002789	4C	seob9635 4C
nucleosome assembly protein 1-like 3	NAPIL3	4675	NP_004529	NM_004538	4C	seob9614 4C
growth hormone inducible transmembrane protein	GHITM	27069	NP_055209	NM_014394	4C	seob9574 4C
ubiquitin-conjugating enzyme E2-like	UEV3	55293	NP_060784	NM_018314	4C	seob9552
FK506 binding protein 14, 22 kDa	FKBP14	55033	NP_060416	NM_017946	4C	seob9543
SAC1 suppressor of actin mutations 1-like (yeast)	SACMIL	22908	NP_054735	NM_014016	4C	seob9485
ubiquitin-conjugating enzyme E2, J1 (UBC6 homolog,	UBE2J1	51465	NP_057105	NM_016021	4C	seob9480 4C
DC2 protein	DC2	58505	NP_067050	NM_021227	4C	seob9430 4C
golgi reassembly stacking protein 2, 55kDa	GORASP2	26003	NP_056345	NM_015530	40	seob9353 4C
leucyl-tRNA synthetase	LARS	51520	NP_064502	NM_020117	4C	seob9282 4C
ubiquitin-conjugating enzyme E2E I (UBC4/5 homolog	UBE2E1	7324	NP_872607	NM_003341	40	seob9266 4C
proteasome (prosome, macropain) subunit, alpha typ	PSMA4	5685	NP_002780	NM_002789	4C	seob9241 4C
proline-rich nuclear receptor coactivator 1	PNRC1	10957	NP_006804	NM_006813	4C	seob9145_4C
multiple coagulation factor deficiency 2	MCFD2	90411	NP_644808	NM_139279	4C	seob9067_4C
general transcription factor IIH, polypeptide 2, 4	GTF2H2	2966	NP_001506	NM_001515	4C	seob8817 4C
				NM_032245	4C	seob8807
zinc finger protein 485	ZNF485	220992	NP_660355	NM_145312	4C	seob8693
WD and tetratricopeptide repeats 1	WDTCI	23038	NP_055838	NM_015023	4C	seob8501
hypothetical protein BC009518	LOC90799	90799	NP_612372	NM_138363	4C	seob8368
chromosome 20 open reading frame 13	C20orf13	55617	NP_060184	NM_017714	4C	seob8355

9872 (4C NM_024537 NP_078813 79587 FL112118 hypothetical protein FL112118 9882 (4C NM_007114 NP_002043 7110 TMFI TATA clement modulatory factor I 9882 (4C) NM_002113 NP_002043 7110 TMFI TATA clement modulatory factor I 9876 (4C) NM_002114 NP_002046 154881 TEXTO testis expressed gene I0 9884 (4C) NM_025146 NP_003297 22988 GNRHR genadomopin-releasing hormone receptor 9894 (4C) NM_025161 NP_078467 280636 SELH selanoprotein H 9895 (4C) NM_018993 NP_06866 54433 RIN2 Rus and Rab interactor 2 9924 (4C) NM_018993 NP_06868 54433 RIN2 Rus and Rab interactor 2 9924 (4C) NM_018903 NP_068788 29980 DONSON downstream neighbor of SON 9924 (4C) NM_024518 NP_068788 116496 Clor24 chromosome 1 open reading frame 13 9925 (4C) NM_002458 NP_003371 7431 VIM </th <th>hypothetical protein BC013035</th> <th>LOC114926</th> <th>114926</th> <th>NP_612445</th> <th>NM_138436</th> <th>4C</th> <th>seoc1948</th>	hypothetical protein BC013035	LOC114926	114926	NP_612445	NM_138436	4C	seoc1948
NM_02437 NP_078813 79587 FLJ12118 NM_007114 NP_009045 7110 TMF1 NM_007113 NP_009045 7110 TMF1 NM_0071746 NP_002104 3078:3079 HFL1 NM_002113 NP_060216 5481 TEX10 NM_002466 NP_079422 80218 MAK3P NM_002466 NP_00397 2798 GNRHR NM_170746 NP_078787 79441 C4orf15 NM_017613 NP_065738 29980 DONSON NM_017613 NP_665738 29980 DONSON NM_052966 NP_443198 116496 C1orf24 NM_0052966 NP_002537 4982 TNFRSF11B NM_003380 NP_003371 7431 VIM NM_004885 NP_003371 7431 VIM NM_004885 NP_079105 79887 FL122662 NM_004888 NP_006991 51056 LAP3 NM_004888 NP_0067248 12709 Ckb	Down syndrome critical region gene 5	DSCR5	51227	NP_710149	NM_016430	†C	seoc1664
NM_024537 NP_078813 79587 FL112118 hypothetical NM_007114 NP_009045 7110 TMF1 TATA eleme NM_007114 NP_009045 7110 TMF1 TATA eleme NM_0071746 NP_002104 3078; 3079 HFL1 H factor (cor NM_002146 NP_060216 54881 TEX10 testis express NM_00406 NP_060397 2798 GNRHR gonadotropit NM_170746 NP_000397 2798 GNRHR gonadotropit NM_018993 NP_061866 54453 RIN2 Ras and Rab NM_018993 NP_065807 79441 C4orf15 chromosome NM_01893 NP_065738 29980 DONSON downstream NM_0145808 NP_065738 29980 DONSON downstream NM_0152966 NP_443198 116496 C1orf24 chromosome NM_002546 NP_002537 4982 TNFRSF11B tumor necros NM_004879 NP_078164 394 ARHGAP5 <td>tyrosine 3-monooxygenase/tryptophan 5-monooxygenas</td> <td>YWHAG</td> <td>7532</td> <td>NP_036611</td> <td>NM_012479</td> <td>‡C</td> <td>seoc1593</td>	tyrosine 3-monooxygenase/tryptophan 5-monooxygenas	YWHAG	7532	NP_036611	NM_012479	‡C	seoc1593
NM_024537 NP_078813 79587 FLJ12118 NM_007114 NP_009045 7110 TMF1 NM_007113 NP_002104 3078; 3079 HFL1 NM_002113 NP_002104 3078; 3079 HFL1 NM_002113 NP_060216 54881 TEX10 NM_00246 NP_079422 80218 MAK3P NM_00246 NP_000397 2798 GNRHR NM_018993 NP_061866 54453 RIN2 NM_017613 NP_078787 79441 C4orf15 NM_017613 NP_665807 136319 MTPN NM_022546 NP_443198 116496 C1orf24 NM_003380 NP_003371 7431 VIM NM_001173 NP_003371 7431 VIM NM_024829 NP_079105 79887 FLJ22662 NM_002794 NP_002785 5690 PSMB2 NM_015907 NP_065991 51056 LAP3 NM_012733 NP_067248 12709 Ckb	ATPase, H+ transporting, lysosomal 13kDa, V1 subun	ATP6V1G1	9550	NP_004879	NM_004888	t)	seoc1539
NM_024537 NP_078813 79587 FLJ12118 NM_007114 NP_009045 7110 TMF1 NM_007114 NP_009045 7110 TMF1 NM_002113 NP_009045 7110 TMF1 NM_002114 3078: 3079 HFL1 NM_002146 NP_060216 54881 TEX10 NM_0025146 NP_079422 80218 MAK3P NM_170746 NP_079422 80218 MAK3P NM_170746 NP_0734467 280636 SELH NM_170746 NP_061866 54453 RIN2 NM_017613 NP_065738 29980 DONSON NM_0145808 NP_665807 136319 MTPN NM_002346 NP_002537 4982 TNFRSF11B NM_003380 NP_003371 7431 VIM NM_144658 NP_003371 7431 VIM NM_024829 NP_079105 79887 FLJ22662 NM_002794 NP_002785 5690 PSMB2 NM_015907 <	creatine kinase, brain	Ckb	12709	NP_067248	NM_021273	ţ)	seoc1264 4C
NIM_024537 NP_078813 79587 FLJ12118 NIM_007114 NP_009045 7110 TMF1 NM_007114 NP_009045 7110 TMF1 NM_002113 NP_002104 3078; 3079 HFL1 NM_0021146 NP_006216 54881 TEX10 NM_0025146 NP_079422 80218 MAK3P NM_170746 NP_00397 2798 GNRHR NM_170746 NP_061866 54453 RIN2 NM_018993 NP_0685787 79441 C4orf15 NM_0145808 NP_665738 29980 DONSON NM_0145808 NP_665807 136319 MTPN NM_0025966 NP_443198 116496 C1orf24 NM_003380 NP_003371 7431 VIM NM_144658 NP_003371 7431 VIM NM_0024829 NP_003371 7431 VIM NM_024558 NP_002785 5690 PSMB2 NM_024558 NP_0078834 79609 C14orf138	leucine aminopeptidase 3	LAP3	51056	NP_056991	NM_015907	ťC T	seoc1234 4C
NM_024537 NP_078813 79587 FLJ12118 NM_007114 NP_009045 7110 TMF1 NM_007114 NP_009045 7110 TMF1 NM_002113 NP_002104 3078; 3079 HFL1 NM_017746 NP_06216 54881 TEX10 NM_000406 NP_079422 80218 MAK3P NM_170746 NP_00397 2798 GNRHR NM_170746 NP_061866 54453 SELH NM_018993 NP_065866 54453 RIN2 NM_017613 NP_665738 29980 DONSON NM_0145808 NP_665807 136319 MTPN NM_002546 NP_443198 116496 Clorf24 NM_003380 NP_002537 4982 TNFRSF11B NM_00173 NP_003371 7431 VIM NM_0044658 NP_653259 139818 DOCK11 NM_002794 NP_002785 5690 PSMB2		C14orf138	79609	NP_078834	NM_024558	†)	seoc1230 4C
NM_024537 NP_078813 79587 FLJ12118 NM_007114 NP_009045 7110 TMF1 NM_002113 NP_002104 3078; 3079 HFL1 NM_017746 NP_060216 54881 TEX10 NM_0177746 NP_079422 80218 MAK3P NM_170746 NP_000397 2798 GNRHR NM_170746 NP_0734467 280636 SELH NM_017613 NP_061866 54453 RIN2 NM_017613 NP_665807 136319 MTPN NM_052966 NP_443198 116496 Clorf24 NM_002546 NP_002537 4982 TNFRSF11B NM_003380 NP_003371 7431 VIM NM_144658 NP_003371 7431 VIM NM_024829 NP_079105 79887 FLJ22662	proteasome (prosome, macropain) subunit, beta type	PSMB2	5690	NP_002785	NM_002794	†C	seoc1203
NM_024537 NP_078813 79587 FLJ12118 NM_007114 NP_009045 7110 TMF1 NM_002113 NP_009045 7110 TMF1 NM_002113 NP_009046 3078; 3079 HFL1 NM_017746 NP_060216 54881 TEX10 NM_0025146 NP_079422 80218 MAK3P NM_170746 NP_000397 2798 GNRHR NM_0170746 NP_0734467 280636 SELH NM_017613 NP_068866 54453 RIN2 NM_017613 NP_665738 29980 DONSON NM_052966 NP_443198 116496 C1orf24 NM_003380 NP_003371 7431 VIM NM_01173 NP_003371 7431 VIM NM_144658 NP_653259 139818 DOCK11	hypothetical protein FLJ22662	FLJ22662	79887	NP_079105	NM_024829	Ċ C	seoc1175
NM_024537 NP_078813 79587 FLJ12118 NM_007114 NP_009045 7110 TMF1 NM_007114 NP_009045 7110 TMF1 NM_002113 NP_009045 7110 TMF1 NM_002113 NP_002104 3078; 3079 HFL1 NM_002146 NP_060216 54881 TEX10 NM_025146 NP_079422 80218 MAK3P NM_170746 NP_000397 2798 GNRHR NM_018993 NP_061866 54453 RIN2 NM_017613 NP_078787 79441 C4orf15 NM_0145808 NP_665738 29980 DONSON NM_0052966 NP_443198 116496 C1orf24 NM_003380 NP_003371 7431 VIM NM_003380 NP_001164 394 ARHGAP5	dedicator of cytokinesis 11	DOCK11	139818	NP_653259	NM_144658	Ť	seoc1159
NM_024537 NP_078813 79587 FLJ12118 NM_007114 NP_009045 7110 TMF1 NM_002113 NP_009045 7110 TMF1 NM_002113 NP_009045 7110 TMF1 NM_002113 NP_0002104 3078; 3079 HFL1 NM_017746 NP_060216 54881 TEX10 NM_00406 NP_079422 80218 MAK3P NM_170746 NP_000397 2798 GNRHR NM_170746 NP_061866 54453 RIN2 NM_018993 NP_061866 54453 RIN2 NM_017613 NP_065738 29980 DONSON NM_052966 NP_443198 116496 C1orf24 NM_002546 NP_002537 4982 TNFRSF11B NM_003380 NP_003371 7431 VIM		ARHGAP5	394	NP_001164	NM_001173	Ť.	seoc1118
NM_024537 NP_078813 79587 FLJ12118 NM_007114 NP_009045 7110 TMF1 NM_007114 NP_009045 7110 TMF1 NM_002113 NP_002104 3078; 3079 HFL1 NM_017746 NP_060216 54881 TEX10 NM_0025146 NP_079422 80218 MAK3P NM_170746 NP_00397 2798 GNRHR NM_018993 NP_061866 54453 RIN2 NM_017613 NP_078787 79441 C4orf15 NM_0145808 NP_665738 29980 DONSON NM_052966 NP_443198 116496 C1orf24 NM_002546 NP_002537 4982 TNFRSF11B	vimentin	MIA	7431	NP_003371	NM_003380	Ţ.	seoc1025
NM_024537 NP_078813 79587 FLJ12118 NM_007114 NP_009045 7110 TMF1 NM_002113 NP_002104 3078; 3079 HFL1 NM_017746 NP_060216 54881 TEX10 NM_025146 NP_079422 80218 MAK3P NM_170746 NP_000397 2798 GNRHR NM_0170746 NP_734467 280636 SELH NM_024511 NP_061866 54453 RIN2 NM_017613 NP_665738 29980 DONSON NM_145808 NP_665807 136319 MTPN NM_052966 NP_443198 116496 Clorf24		TNFRSF11B	4982	NP_002537	NM_002546	÷.	seoc1023
NM_024537 NP_078813 79587 FLJ12118 NM_007114 NP_009045 7110 TMF1 NM_002113 NP_002104 3078; 3079 HFL1 NM_017746 NP_060216 54881 TEX10 NM_025146 NP_079422 80218 MAK3P NM_000406 NP_000397 2798 GNRHR NM_170746 NP_734467 280636 SELH NM_018993 NP_061866 54453 RIN2 NM_024511 NP_078787 79441 C4orf15 NM_145808 NP_665738 29980 DONSON		Clorf24	116496	NP_443198	NM_052966	t)	seoc0999
NM_024537 NP_078813 79587 FLJ12118 NM_007114 NP_009045 7110 TMF1 NM_002113 NP_002104 3078; 3079 HFL1 NM_017746 NP_060216 54881 TEX10 NM_025146 NP_079422 80218 MAK3P NM_00406 NP_000397 2798 GNRHR NM_170746 NP_734467 280636 SELH NM_018993 NP_061866 54453 RIN2 NM_024511 NP_078787 79441 C4orf15 NM_017613 NP_665738 29980 DONSON	myotrophin	MTPN	136319	NP_665807	NM_145808	ţ,	seoc0924 4C
NM_024537 NP_078813 79587 FLJ12118 NM_007114 NP_009045 7110 TMF1 NM_002113 NP_002104 3078; 3079 HFL1 NM_017746 NP_060216 54881 TEX10 NM_025146 NP_079422 80218 MAK3P NM_000406 NP_000397 2798 GNRHR NM_170746 NP_734467 280636 SELH NM_018993 NP_061866 54453 RIN2 NM_024511 NP_078787 79441 C4orf15	downstream neighbor of SON	DONSON	29980	NP_665738	NM_017613	t)	seoc0866 4C
NM_024537 NP_078813 79587 FLJ12118 NM_007114 NP_009045 7110 TMF1 NM_002113 NP_002104 3078; 3079 HFL1 NM_017746 NP_060216 54881 TEX10 NM_025146 NP_079422 80218 MAK3P NM_000406 NP_000397 2798 GNRHR NM_170746 NP_734467 280636 SELH NM_018993 NP_061866 54453 RIN2		C4orf15	79441	NP_078787	NM_024511	ţ	seoc0778 4C
NM_024537 NP_078813 79587 FLJ12118 NM_007114 NP_009045 7110 TMF1 NM_002113 NP_002104 3078; 3079 HFL1 NM_017746 NP_060216 54881 TEX10 NM_025146 NP_079422 80218 MAK3P NM_170746 NP_734467 280636 SELH	Ras and Rab interactor 2	RIN2	54453	NP_061866	NM_018993	tC .	seoc0775 4C
NM_024537 NP_078813 79587 FLJ12118 NM_007114 NP_009045 7110 TMF1 NM_002113 NP_002104 3078; 3079 HFL1 NM_017746 NP_060216 54881 TEX10 NM_025146 NP_079422 80218 MAK3P NM_000406 NP_000397 2798 GNRHR		SELH		NP_734467	NM_170746	t)	seoc0513 4C
NM_024537 NP_078813 79587 FLJ12118 NM_007114 NP_009045 7110 TMF1 NM_002113 NP_002104 3078; 3079 HFL1 NM_017746 NP_060216 54881 TEX10 NM_025146 NP_079422 80218 MAK3P	gonadotropin-releasing hormone receptor	GNRHR	2798	NP_000397	NM_000406	C	seoc0394 4C
NM_024537 NP_078813 79587 FLJ12118 NM_007114 NP_009045 7110 TMF1 NM_002113 NP_002104 3078; 3079 HFL1 NM_017746 NP_060216 54881 TEX10	likely ortholog of mouse Mak3p homolog (S. cerevis	МАКЗР	80218	NP_079422	NM_025146	7	seoc0369 /
NM_024537 NP_078813 79587 FLJ12118 NM_007114 NP_009045 7110 TMF1 NM_002113 NP_002104 3078; 3079 HFL1	testis expressed gene 10	TEX10	54881	NP_060216	NM_017746	7	seoc0276 4C
NM_024537 NP_078813 79587 FLJ12118 NM_007114 NP_009045 7110 TMF1	H factor (complement)-like 1	HFLI	3078; 3079	NP_002104	NM_002113	E O	seob9946 4C
NM_024537	TATA element modulatory factor 1	TMFI	7110	NP_009045	NM_007114	.C	seob9882 -
		FLJ12118	79587	NP_078813	NM_024537	Ŋ	seob9872 4C

4C NM_144781 NP_659005 5134 PDCD2 4C NM_133325 NP_579866 55274 PHF10 4C NM_0020948 NP_065999 57708 MI-ER1 4C NM_005204 NP_005195 1326 MAP3K8 4C NM_001207 NP_001198 689; 220793 BTF3 4C NM_017333 NP_060203 54872 FLJ20265 4C NM_017733 NP_060204 4627 MYH9 4C NM_017734 NP_060204 54873 PALMD 4C NM_0173060 NP_0775085 831 CAST 4C NM_0173060 NP_061932 54543 TOMM7 4C NM_0167549 NP_066927 6171 RPL41 4C NM_002160 NP_002151 3371 TNC 4C NM_003877 NP_066927 6171 RPL41 4C NM_003877 NP_069911 55825 PECR 4C NM_003876 NP_069911 5	leucine-rich repeats and immunoglobulin-like domai	LRIG1	26018	NP_056356	NM_015541	4C	seoc5285 4C
4C NM_144781 NP_659005 5134 PDCD2 programmed of tenascin C (he case) 4C NM_133325 NP_579866 55274 PHF10 PHD finger programmed of tenascin C (he case) 4C NM_020948 NP_065999 57708 M1-ER1 mesoderm ind 4C NM_005204 NP_005195 1326 MAP3K8 mitogen-activ 4C NM_01207 NP_001198 689; 220793 BTF3 basic transcrit 4C NM_017733 NP_060203 54872 FLJ20265 hypothetical p 4C NM_017734 NP_060204 4627 MYH9 myosin, heavy 4C NM_0173060 NP_775085 831 CAST calpastatin 4C NM_019059 NP_061932 54543 TOMM7 transcin C (he 4C NM_002160 NP_002151 3371 TNC tenascin C (he 4C NM_002104 NP_006927 6171 RPL41 ribosomal pro 4C NM_002056 NP_0066927 6171 PCBP2	GM2 ganglioside activator protein	GM2A	2760; 401213	NP_000396	NM_000405	Ć	seoc4960 4C
4C NM_144781 NP_659005 5134 PDCD2 programmed of	tetratricopeptide repeat domain 5	TTC5	91875	NP_612385	NM_138376	†)	seoc4909
4C NM_144781 NP_659005 5134 PDCD2 programmed of	peroxisomal trans-2-enoyl-CoA reductase	PECR	55825	NP_060911	NM_018441	t)	seoc4505
4C NM_144781 NP_659005 5134 PDCD2 programmed of the programme of the programmed of t	poly(rC) binding protein 2	РСВР2	5094	NP_114366	NM_005016	tc	seoc4416
4C NM_144781 NP_659005 5134 PDCD2 programmed of tenascin C (he case) 4C NM_133325 NP_659005 5134 PDCD2 programmed of tenascin C (he case) 4C NM_020948 NP_065999 57708 M1-ER1 mesoderm ind 4C NM_005204 NP_065999 57708 M1-ER1 mesoderm ind 4C NM_001207 NP_065999 57708 M1-ER1 mesoderm ind 4C NM_001207 NP_005195 1326 MAP3K8 mitogen-activ 4C NM_001207 NP_0060203 54872 FLJ20265 hypothetical p 4C NM_0017733 NP_060203 54873 PALMD palmdelphin 4C NM_1173060 NP_775085 831 CAST calpastatin 4C NM_002160 NP_061932 54543 TOMM7 translocase of 4C NM_002160 NP_002151 3371 TNC tenascin C (he 4C NM_002160 NP_066927 6171 RPL41		CTNNBLI	56259	NP_110517	NM_030877	1 C	seoc4380
4C NM_144781 NP_659005 5134 PDCD2 programmed of tenascin C (he documents) 4C NM_133325 NP_579866 55274 PHF10 PHD finger programmed of tenascrity 4C NM_020948 NP_065999 57708 M1-ER1 mesoderm ind 4C NM_005204 NP_005195 1326 MAP3K8 mitogen-active 4C NM_001207 NP_001198 689; 220793 BTF3 basic transcrity 4C NM_0017733 NP_060203 54872 FLJ20265 hypothetical programmed of transcrity 4C NM_002473 NP_060203 54872 FLJ20265 hypothetical programmed of transcrity 4C NM_017734 NP_060203 54872 FLJ20265 hypothetical programmed of transcrity 4C NM_0177360 NP_775085 831 CAST calpastatin 4C NM_002160 NP_061932 54543 TOMM7 translocase of transcrity 4C NM_002160 NP_002151 3371 TNC tenascin C (he 4C	protocadherin beta 16	PCDHB16	57717	NP_066008	NM_020957	10	seoc4288
4C NM_144781 NP_659005 5134 PDCD2 programmed of tenascin C (he 4C NM_133325 NP_579866 55274 PHF10 PHD finger pn 4C NM_020948 NP_065999 57708 M1-ER1 mesoderm ind 4C NM_005204 NP_005195 1326 MAP3K8 mitogen-activ 4C NM_001207 NP_001198 689: 220793 BTF3 basic transcrip 4C NM_017733 NP_060203 54872 FLJ20265 hypothetical p 4C NM_017734 NP_060204 4627 MYH9 myosin, heavy 4C NM_0173060 NP_775085 831 CAST calpastatin 4C NM_019059 NP_061932 54543 TOMM7 translocase of 4C NM_002160 NP_002151 3371 TNC tenascin C (he 4C NM_002160 NP_006927 6171 RPL41 ribosomal pro	spermine synthase	SMS	6611	NP_004586	NM_004595	t)	seoc4187
4C NM_144781 NP_659005 5134 PDCD2 programmed of tenascin C (he 4C NM_133325 NP_579866 55274 PHF10 PHD finger pn 4C NM_020948 NP_065999 57708 MI-ERI mesoderm ind 4C NM_005204 NP_005195 1326 MAP3K8 mitogen-activ 4C NM_001207 NP_001198 689; 220793 BTF3 basic transcrip 4C NM_017733 NP_060203 54872 FLJ20265 hypothetical p 4C NM_017734 NP_060204 54873 PALMD palmdelphin 4C NM_019059 NP_061932 54543 TOMM7 translocase of 4C NM_002160 NP_002151 3371 TNC tenascin C (he	ribosomal protein L41	RPL41	6171	NP_066927	NM_021104	t)	seoc4052
4C NM_144781 NP_659005 \$134 PDCD2 programmed of	tenascin C (hexabrachion)	TNC	3371	NP_002151	NM_002160	†C	seoc3993
4C NM_144781 NP_659005 5134 PDCD2 programmed of					XM_167549	Ċ.	seoc3980 4C
4C NM_144781 NP_659005 5134 PDCD2 programmed of	tenascin C (hexabrachion)	TNC	3371	NP_002151	NM_002160	ţ.	seoc3876
4C NM_144781 NP_659005 5134 PDCD2 programmed cell death 2 4C NM_133325 NP_579866 55274 PHF10 PHD finger protein 10 4C NM_020948 NP_065999 57708 MI-ER1 mesoderm induction early response 1 4C NM_005204 NP_005195 1326 MAP3K8 mitogen-activated protein kinase kinase kinase kinase kinase kinase kinase 4C NM_001207 NP_001198 689; 220793 BTF3 basic transcription factor 3 4C NM_017733 NP_060203 54872 FLJ20265 hypothetical protein FLJ20265 4C NM_002473 NP_060204 4627 MYH9 myosin, heavy polypeptide 9, non-muscle 4C NM_0173060 NP_775085 831 CAST calpastatin	translocase of outer mitochondrial membrane 7 homo	ТОММ7	54543	NP_061932	NM_019059	tC	seoc3603
4C NM_144781 NP_659005 5134 PDCD2 programmed cell death 2 4C NM_133325 NP_579866 55274 PHF10 PHD finger protein 10 4C NM_020948 NP_065999 57708 MI-ER1 mesoderm induction early response 1 4C NM_005204 NP_005195 1326 MAP3K8 mitogen-activated protein kinase kina	calpastatin	CAST	831	NP_775085	NM_173060	†C	seoc3588
4C NM_144781 NP_659005 5134 PDCD2 programmed cell death 2 4C NM_133325 NP_579866 55274 PHF10 PHD finger protein 10 4C NM_020948 NP_065999 57708 MI-ER1 mesoderm induction early response 1 4C NM_005204 NP_005195 1326 MAP3K8 mitogen-activated protein kinase kina	palmdelphin	PALMD	54873	NP_060204	NM_017734	ţ.	seoc2824
4C NM_144781 NP_659005 5134 PDCD2 programmed cell death 2 4C NM_133325 NP_579866 55274 PHF10 PHD finger protein 10 4C NM_020948 NP_065999 57708 M1-ER1 mesoderm induction early response 1 4C NM_005204 NP_005195 1326 MAP3K8 mitogen-activated protein kinase kina	myosin, heavy polypeptide 9, non-muscle	МҮН9	4627	NP_002464	NM_002473	ţ.	seoc2447
4C NM_144781 NP_659005 5134 PDCD2 programmed cell death 2 4C NM_133325 NP_579866 55274 PHF10 PHD finger protein 10 4C NM_020948 NP_065999 57708 MI-ER1 mesoderm induction early response 1 4C NM_005204 NP_005195 1326 MAP3K8 mitogen-activated protein kinase kinase kinase kinase 4C XM_210358 NP_001198 689; 220793 BTF3 basic transcription factor 3 4C NM_020555 NP_001198 689; 220793 BTF3 basic transcription factor 3	hypothetical protein FLJ20265	FLJ20265	54872	NP_060203	NM_017733	†C	seoc2264
4C NM_144781 NP_659005 5134 PDCD2 programmed cell death 2 4C NM_133325 NP_579866 55274 PHF10 PHD finger protein 10 4C NM_020948 NP_065999 57708 MI-ER1 mesoderm induction early response 1 4C NM_005204 NP_005195 1326 MAP3K8 mitogen-activated protein kinase kina					NM_020555	ť	seoc2253
4C NM_144781 NP_659005 5134 PDCD2 programmed cell death 2 4C NM_133325 NP_579866 55274 PHF10 PHD finger protein 10 4C NM_020948 NP_065999 57708 MI-ER1 mesoderm induction early response 1 4C NM_005204 NP_005195 1326 MAP3K8 mitogen-activated protein kinase kinase kinase 4C XM_210358 March 12 March 13 March 13 March 14 March 16 M	basic transcription factor 3	BTF3	689; 220793	NP_001198	NM_001207	†C	seoc2246
4C NM_144781 NP_659005 5134 PDCD2 programmed cell death 2 4C NM_133325 NP_579866 55274 PHF10 PHD finger protein 10 4C NM_020948 NP_065999 57708 MI-ER1 mesoderm induction early response 1 4C NM_005204 NP_005195 1326 MAP3K8 mitogen-activated protein kinase kinase kinase					XM_210358	t	seoc2226
4C NM_144781 NP_659005 5134 PDCD2 4C NM_133325 NP_579866 55274 PHF10 4C NM_020948 NP_065999 57708 M1-ER1	mitogen-activated protein kinase kinase kinase 8	MAP3K8	1326	NP_005195	NM_005204	†C	seoc2191
4C NM_144781 NP_659005 5134 PDCD2 4C NM_133325 NP_579866 55274 PHF10	mesoderm induction early response 1	MI-ERI	57708	NP_065999	NM_020948	tC	seoc2173
4C NM_144781 NP_659005 5134 PDCD2	PHD finger protein 10	PHF10	55274	NP_579866	NM_133325	7	seoc2131
		PDCD2	5134	NP_659005	NM_144781	t)	seoc2029
4C NM 017751 NP 060421 55627 FLJ20297	hypothetical protein FLJ20297	FLJ20297	55627	NP_060421	NM_017751	TC	seoc1996

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	NM_024692		INM 002356	The state of the s	NM 003253	61/400 TAIA;	NM 004710
	NP 078968	146700	NP 002347	111_005244	ND 003744	NP_004710	
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Typothetical protein FLJ21069	The state of the s	myristoylated alanine-rich protein kingse C subst-	Gleptom and memorals	T-cell lymphoma invasion and metastacic i	' s' "" " s''' " s' line-rien 2, interacti	Splicing factor argining/caring	